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SEARCH REQUEST FORM

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23

Requester's Full Name: ANN LAM Examiner #: 77568 Date: 8/26/03  
 Art Unit 1641 Phone Number 306-5560 Serial Number 09/935417  
 Mail Box and Bldg/Room Location: 7E12 Results Format Preferred (circle) PAPER DISK E-MAIL

If more than one search is submitted, please prioritize searches in order of need.

Please provide a detailed statement of the search topic, and describe as specifically as possible the subject matter to be searched. Include the elected species or structures, keywords, synonyms, acronyms, and registry numbers, and combine with the concept or utility of the invention. Define any terms that may have a special meaning. Give examples or relevant citations, authors, etc. if known. Please attach a copy of the cover sheet, pertinent claims, and abstract.

Title of Invention: Composite Expandable Device w/ Polymeric Covering + bioactive coating...  
 Inventors (please provide full names): Leon V. Rudakov & Mir A. Imrhan  
+ Linh Dinh; Ara Davidian & ~~Kevin~~ Kevin T. Larkin  
 Earliest Priority Filing Date: Aug 30, 1999

\*For Sequence Searches Only\* Please include all pertinent information (parent, child, divisional, or issued patent numbers) along with the appropriate serial number.

Please search peptide that includes the  
 amino acid sequence ID No: 1 (claim 19)  
 (peptide enhances endothelial cell growth)  
 Thank you!  
 Ann

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AA  
 1-15

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OM protein - protein search, using sw model

Run on: August 29, 2003, 17:49:44 ; Search time 83 Seconds

28.686 Million cell updates/sec

Title: US-09-935-417-1

Sequence: 1 GTPGPQGIAGQRGV 15

Scoring table: BLOSUM62

Searched: 1107863 seqs, 158726573 residues

Total number of hits satisfying chosen parameters: 1107863

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Minimum DB seq length: 0
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Maximum DB seq Length: 20000000000

Post-processing:	Minimum Match	0%
	Maximum Match	100%

Listing first 45 summaries

Database : A\_Geneseq\_19Jun03:\*

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22:	/SDSI/gcgdata/geneseq/geneseqp-emb1/AA2001.DAT *
23:	/SDSI/gcgdata/geneseq/geneseqp-emb1/AA2002.DAT *
24:	/SDSI/gcgdata/geneseq/geneseqp-emb1/AA2003.DAT *

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	80	100.0	15	12	AAK11114	Collagen peptide a
2	80	100.0	15	14	AAK38476	Sequence of peptid
3	80	100.0	15	18	AAW27491	Cell binding peptid
4	80	100.0	15	18	AAW16825	Collagen binding p
5	80	100.0	15	20	AAV29991	collagen cell bind
6	80	100.0	15	20	AAV29587	collagen fibronect
7	80	100.0	15	22	AAAG67402	Synthetic peptide
8	80	100.0	15	23	ABP51951	Portion of an al c
9	80	100.0	15	23	ABB10111	Collagen cell bind

10	80	100.0	16	17	AAR92859	Collagen fragment
11	80	100.0	16	21	AAR76868	Collagen receptor
12	80	100.0	19	22	AAB35632	Collagenase cleave
13	80	100.0	25	20	AAV07306	Collagen assembly
14	80	100.0	134	22	ABU53024	Human testes-deri
15	80	100.0	144	22	ABU53007	Human testes-deri
16	80	100.0	144	22	ABU53008	Human testes-deri
17	80	100.0	333	22	AAE02713	Recombinant human
18	80	100.0	333	22	AAB68067	Recombinant human
19	80	100.0	416	22	AAE02711	Human alpha (1)
20	80	100.0	416	22	AAB68065	Human alpha (1)
21	80	100.0	500	22	AAE02708	Human alpha (1)
22	80	100.0	500	22	AAB68062	Amino acid sequen
23	80	100.0	510	22	AAE02712	Recombinant human
24	80	100.0	510	22	AAB68066	Amino acid sequen
25	80	100.0	662	22	AAE02718	Human alpha (1)
26	80	100.0	662	22	AAB68072	Amino acid sequen
27	80	100.0	822	22	AAV06240	Mouse recombinant
28	80	100.0	936	22	AAE070107	Gelatin protein.
29	80	100.0	1057	21	AAV84541	Amino acid sequen
30	80	100.0	1057	21	AAV84544	A human collagen
31	80	100.0	1058	21	AAV84403	Amino acid sequen
32	80	100.0	1107	17	AAR89472	Collagen/decorin(
33	80	100.0	1107	21	AAV84540	Amino acid sequen
34	80	100.0	1169	17	AAR89469	Collagen/BMP-2B f
35	80	100.0	1169	21	AAV84537	Amino acid sequen
36	80	100.0	1171	17	AAR89470	Collagen/TGF-beta
37	80	100.0	1171	21	AAV84538	A chimeric collag
38	80	100.0	1341	16	AAV71701	Collagen alpha 1
39	80	100.0	1341	21	AAV96122	Collagen type I-a
40	80	100.0	1341	23	ABB80733	Collagen type I-a
41	80	100.0	1341	23	ABB09625	Amino acid sequen
42	80	100.0	1341	23	AAE16475	Human collagen al
43	80	100.0	1388	17	AAR89471	Collagen/decorin
44	80	100.0	1388	21	AAV84539	Amino acid sequen
45	80	100.0	1411	21	AAV56800	Human preproalpha

## ALIGNMENTS

```

RESULT 1
AAR11114
ID  AAR11114 standard; peptide; 15 AA

```

AC AAR11114;

DT	25-MAR-2003	(updated)
DT	17-MAY-1991	(first entry)

Collagen peptide analogue.

Collagen alpha-1 chain; cell adhesion; vertebrates

OS synthetic.

PN W09102537-A.

PD 07-MAR-1991.

PF 13-AUG-1990; 90WO-US04538.  
YY

PR 14-AUG-1989; 89US-0393621.  
VY

PA (REGC ) UNIV CALIFORNIA.  
YY

PI Bhatnagar RS;  
YY

DR WPI; 1991-087110/12  
XY

PT Synthetic peptide(s) analogous to collagen - promote cell adhesion

PS Claim 1; page 16; 20pp; English

XX This peptide corresponds to a region of the alpha-1 chain of collagen.  
 CC It is useful in a compn. for promoting vertebrate cell (esp.  
 CC fibroblast) adhesion to a substrate. It is free from natural  
 CC folding, glycosylation, cross-linking, hydroxylation and association  
 CC with other peptide chains.  
 CC (Updated on 25-MAR-2003 to correct PA field.)

XX Sequence 15 AA:

Query Match 100.0%; Score 80; DB 12; Length 15;  
 Best Local Similarity 100.0%; Pred. NO. 6.9e-05;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCVV 15  
 Db 1 GTPGPGIAGGRCVV 15

RESULT 2  
 AAR38476 standard; peptide; 15 AA.

AC AAR38476;  
 DT 25-MAR-2003 (updated)  
 DT 02-DEC-1993 (first entry)

DE Sequence of peptide P-15 which spans approx. residues 766-780 of the  
 alpha-1(I) chain of collagen.

XX Synthetic peptide; alpha-1(I) chain; collagen; binding; P-15.

OS Synthetic.

PN M09311781-A1.

PD 24-JUN-1993.

PF 03-DEC-1992; 92MO-US10420.

PR 09-DEC-1991; 91US-0804782.

PA (REGC ) UNIV CALIFORNIA.

PI Bhatnagar RS;

DR WPI: 1993-213814/26.

XX Synthetic peptide mimicking collagen binding to cells - used in  
 PT composite with bio-material matrix for soft and hard tissue  
 PT repair or reconstruction

PS Disclosure; Table 1, page 9; 26pp; English.

XX The P-15 peptide spans approx. residues 766-780 of the alpha-1(I)  
 CC chain of collagen. The P-15 region does not occur as a natural  
 CC fragment of collagen nor is it a product of natural enzymatic  
 CC cleavage. The P-15 region represents half of one turn of the collagen  
 CC triple helix. The sequence contd. in P-15 can acquire a conformation  
 CC dramatically different from the triple helical conformation  
 CC generally observed in the rest of the collagen molecule. AAR38477-82  
 CC is a family of synthetic peptide fragments of P-15. They mimic the  
 CC cell binding domain of collagen. The domain includes a core  
 CC sequence that, at physiologic conditions, is folded in a beta-bend  
 CC formed at the Ile-Ala.  
 CC (Updated on 25-MAR-2003 to correct PN field.)

XX Sequence 15 AA:

Query Match 100.0%; Score 80; DB 14; Length 15;  
 Best Local Similarity 100.0%; Pred. NO. 6.9e-05;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCVV 15  
 Db 1 GTPGPGIAGGRCVV 15

RESULT 3  
 AAM27491 standard; peptide; 15 AA.

AC AAM27491;

DT 20-APR-1998 (first entry)

DE Cell binding peptide #1 derived from collagen.

XX Bioreactor; packing material; cell culture; collagen alpha1(I) chain;  
 KW cell binding peptide; matrix.

XX Synthetic.  
 OS Mammalia.

PN US5674848-A.

PD 07-OCT-1997.

PF 03-AUG-1994; 94US-0285570.

PR 14-AUG-1989; 89US-0393621.

PR 09-DEC-1991; 91US-0804782.

PA (REGC ) UNIV CALIFORNIA.

PI Bhatnagar RS;

DR WPI: 1997-502373/46.

XX Bioreactor packing material for cell culture - comprising matrix  
 PT coated with cell binding peptide

PS Claim 1; Col 18; 13pp; English.

XX The present peptide sequence corresponds to a region of the alpha1(I)  
 CC chain of collagen which is sometimes referred to as "P-15". It can be  
 CC used as a cell binding peptide in a new packing material, which is useful  
 CC for cell culture in a bioreactor. The material comprises a matrix formed  
 CC of a biomaterial, i.e. a material that is biologically compatible for in  
 CC vivo applications and for cell culture in vitro, and the cell binding  
 CC peptide. A bioreactor containing the packing material can be used to  
 CC culture cells, e.g. mammalian cells for the production of monoclonal  
 CC antibodies. The peptides are more effective than collagen in promoting  
 CC cell attachment.

XX Sequence 15 AA:

Query Match 100.0%; Score 80; DB 18; Length 15;  
 Best Local Similarity 100.0%; Pred. NO. 6.9e-05;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCVV 15  
 Db 1 GTPGPGIAGGRCVV 15

RESULT 4  
 AAM18825 standard; peptide; 15 AA.

AC AAM18825;

DT 25-MAR-2003 (updated)  
 DT 05-JAN-1998 (first entry)

XX

DE Collagen binding peptide mimic 1.

KW implant; biomaterial matrix; enhanced cell binding; collagen;

KM beta-bend; fold; substrate; reconstructive surgery; bone; ligament;

KW repair; tooth.

XX Synthetic.

OS

XX US5635482-A ★

XX 03-JUN-1997.

XX

XX 22-JUL-1994; 94US-0278878.

XX PF

XX PR 22-JUL-1994; 94US-0278878.

XX PR 14-AUG-1989; 89US-0393621.

XX PR 09-DEC-1991; 91US-0804782.

XX PA (REGC ) UNIV CALIFORNIA.

XX PI Bhatnagar RS;

XX WP; 1997-309859/28.

XX DR

XX PT Implant bearing cell-binding collagen-mimetic peptide - for

XX PP promoting cell attachment

XX PS Claim 1; Column 18; 12pp; English.

XX CC New implants comprise a biomaterial matrix and a peptide carried by the

CC CC matrix, where the peptide has enhanced cell binding with respect to

CC CC collagen and has a domain that mimics collagen binding to cells, the

CC CC domain including at least -Ile-Ala- folded in a beta-bend at

CC CC physiological conditions. The peptide is one of AAW18825-34 or one of 3

CC CC tripeptides (Nac-Ile-Ala-Ala; Ile-Ala-beta Ala; and Nac-Ile-Ala-N-Me).

CC CC The implant is used as a substrate for growing cells, e.g. for use in

CC CC reconstructive surgery, e.g. for bone or ligament repair or as tooth

CC CC implants. The peptide promotes cell attachment to the matrix and also

CC CC cell migration into the matrix when the matrix is porous.

CC CC (Updated on 25-MAR-2003 to correct pf field.)

XX SQ Sequence 15 AA:

OY Query Match Best Local Similarity 100.0%; Score 80; DB 18; Length 15;

Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

DB 1 GTPGPQGIAGORGVV 15  
1 GTPGPQGIAGORGVV 15

RESULT 5

AY29991 standard; peptide; 15 AA.

AAZ29991;

02-DEC-1999 (first entry)

Collagen cell binding domain mimotope #1.

Collagen; cell binding domain; biomaterial; soft tissue repair;  
hard tissue repair; reconstruction; cell surface receptor;  
fibronectin; beta-bend; cartilage; tendon; ligament; bone.

Synthetic.

US5958428-A ★

28-SEP-1999.

97US-0859610.

22-JUL-1994: 94US-0278878.  
PR 14-AUG-1989: 89US-0393621.  
PR 09-DEC-1991: 91US-0804782.  
PA (REGC ) UNIV CALIFORNIA.  
PI Bhatnagar RS;  
DR WPI: 1999-561009/47.  
PT Synthetic peptide additives with enhanced collagen binding affinities  
PT useful for the production of apparatus for soft tissue, cartilage and  
PT bone repair -  
PS Claim 3; Column 25; 16pp; English.  
XX  
XX The present invention describes synthetic peptide additives (SPAs) with  
CC enhanced collagen binding affinities. AAY2991 to AAY3000 represent  
CC specifically claimed examples of the SPA's. The additives comprise  
CC domains that mimic the binding sites of collagen to cells (but with  
CC higher affinity) and promote cell attachment when the additives are  
CC carried on repair or reconstructive apparatus. The SPA may be used in  
CC the construction of apparatus for soft tissue, cartilage, tendon,  
CC ligament and bone repair. The SPA mimics and enhances the binding of  
CC cells to the tissue repair apparatus.  
XX  
XX Sequence 15 AA:  
SQ  
Query Match 100.0%; Score 80; DB 20; Length 15;  
Best Local Similarity 100.0%; Pred. NO. 6.9e-05; Indels 0; Gaps 0  
Matches 15; Conservative 0; Mismatches 0;  
QY 1 GTPGPGGIAGGQGVV 15  
Db 1 GTPGPGGIAGGQGVV 15  
1 GTPGPGGIAGGQGVV 15  
RESULT 6  
AAY29587  
ID AAY29587 standard; peptide; 15 AA.  
XX  
XX AAY29587;  
AC  
XX 18-OCT-1999 (first entry)  
DT  
XX Collagen fibronectin binding region oligopeptide.  
DE  
XX Collagen; fibronectin binding region; tissue regeneration; implant;  
KW internal wound site; biodegradable microparticle.  
KM  
XX Unidentified.  
OS  
XX W09933447-A2.  
XX  
XX 08-JUL-1999.  
PD  
XX 24-DEC-1998; 98WO-US27596.  
PF  
XX 30-DEC-1997; 97US-0000638.  
PR  
XX (MASI ) MASSACHUSETTS INST TECHNOLOGY.  
PA  
XX Yamas IV;  
PI  
XX WPI: 1999-493795/41.  
XX  
XX Biodegradable microparticles for tissue regeneration at an internal  
PT wound site  
PS  
XX Disclosure; Page 8; 25pp; English.  
XX  
XX The present invention describes a porous biodegradable microparticle (1)

CC for tissue regeneration at an internal wound site in a subject. The  
 CC pores of (I) have a diameter 1-300 nm m; (I) has a minimum water content  
 CC of at least about 80%, a minimum specific surface area of at least about  
 CC 103 mm<sup>2</sup> per cm<sup>3</sup> and a diameter 10-1000 micro m. between about 20-80% by  
 CC weight of (I) is biodegraded at the wound site during the time period  
 CC required for a wound of about the same severity, size and tissue type to  
 CC complete about one half of the contraction which normally takes place in  
 CC the absence of (I); and (II) comprises: (i) a three dimensional network  
 CC of polymers which is substantially insoluble under physiological  
 CC conditions; and (ii) one or more specific cell-binding fragments.  
 CC Methods using (I) may be used to treat internal injuries caused to  
 CC internal organs by disease or trauma, and to inhibit wound contraction  
 CC and scar formation. The methods work by preventing contractile cells in  
 CC the vicinity of a wound site (accidentally or surgically induced) on an  
 CC internal organ from inducing contraction at the lesion site. The tissue  
 CC regeneration methods greatly improve the clinical outcomes of patients  
 CC with internal organ and tissue injuries. The present sequence represents  
 CC a collagen fibronectin binding region oligopeptide which is used in as  
 CC part of an example of a specific cell binding fraction which is included  
 CC in a 3-dimensional network of the regeneration template from the present  
 CC invention.

CC Sequence 15 AA:

Query Match 100.0%; Score 80; DB 20; Length 15;  
 Best Local Similarity 100.0%; Pred. No. 6.9e-05;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 GTPPGGAGGRCVV 15  
 |||||  
 DB 1 GTPPGGAGGRCVV 15

RESULT 2

AA67402  
 ID AA67402 standard; peptide: 15 AA.

AC AA67402;

DT 13-NOV-2001 (first entry)

DE Synthetic peptide mimicking cell binding domain of collagen.

KW Cell binding; collagen; cell migration; collagen receptor; tissue repair;  
 KW metalloproteinase; prolyl hydroxylase; tissue reconstruction; arthritis;  
 KW bone repair; tooth implant; ligament repair; scar tissue; osteoporosis;  
 KW bone disease; cartilage repair; joint disease; tendon repair.

OS Synthetic.

PN US6268348-B1.

PD 31-JUL-2001.

PE 08-JUN-1999; 99US-0328347.

PR 22-JUL-1994; 94US-0278878.

PR 20-MAY-1997; 97US-0859610.

PR 14-AUG-1989; 89US-0393621.

PR 09-DEC-1991; 91US-0804782.

PA (REGC) UNIV CALIFORNIA.

PI Bhattachar RS;

DR WPI: 2001-540321/60.

PT New collagen binding synthetic peptide useful for soft and hard tissue  
 PT repair e.g. bone repairs comprises a family of amino acid sequence -

PS Claim 1; Column 25; 16pp; English.  
 XX The present sequence represents a synthetic peptide, which mimics the

CC cell binding domain of collagen. The cell binding ability of the  
 CC peptide is enhanced with respect to collagen. The peptide promotes cell  
 CC migration into porous lattices; binds to collagen receptors; induces  
 CC metalloproteinases; can down regulate prolyl hydroxylase and collagen;  
 CC inhibits cell binding to collagen or inhibits cell migration in vitro.  
 CC The peptide is used for soft and hard tissue repair or reconstruction;  
 CC e.g. bone repair, tooth implants and ligament repair for in vitro uses;  
 CC as an inhibitor of collagen synthesis to block formation of scar tissue  
 CC and thus promotes scarless healing; as bone filling/fusion for  
 CC osteoporosis and other bone diseases; cartilage repair for arthritis and  
 CC other joint disease and tendon repair; for soft tissue repair e.g. nerve,  
 CC organ, skin, vascular, muscle and ophthalmic applications.

CC Sequence 15 AA:

Query Match 100.0%; Score 80; DB 22; Length 15;  
 Best Local Similarity 100.0%; Pred. No. 6.9e-05;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 GTPPGGAGGRCVV 15  
 |||||  
 DB 1 GTPPGGAGGRCVV 15

RESULT 8

ID ABP51951 standard; peptide: 15 AA.

AC ABP51951;

DT 08-OCT-2002 (first entry)

DE Portion of an al chain or collagen peptide sequence SEQ ID NO:1.

KW Delivery; blood; collagen; occlusion; blood vessel; saphenous vein graft.

OS Synthetic.

PN US2002062145-A1.

PD 23-MAY-2002.

PE 22-AUG-2001; 2001US-0935417.

PR 30-AUG-1999; 99US-0385691.

PA (CARD-) CARDIOVASC INC.

PI Rudakov, LV Imran MA, Dinh L, Davidian A, Larkin KT;

DR WPI: 2002-582305/62.

PT Composite expandable device for treating occlusions in blood vessels,  
 PT e.g., saphenous vein grafts, comprises polymeric covering and bioactive  
 PT coating

PS Disclosure; Page 4; 10pp; English.

CC The present invention describes a composite expandable device for  
 CC delivery into a blood vessel comprising an expandable support frame,  
 CC an impervious polymer sleeve extending over the support frame, and a  
 CC coating disposed on the inner and outer surfaces of the polymer sleeve  
 CC for enhancing endothelial cell growth on the polymer sleeve. Also  
 CC described is delivery apparatus for an expandable device comprising a  
 CC shaft, and a balloon mounted on the shaft, where the shaft has a lumen  
 CC for inflating and deflating the balloon. The balloon is formed with  
 CC intermediate portions adapted to receive the expandable device, and  
 CC radiopaque markers carried within the balloon and sized to prevent from  
 CC being dislodged during deployment by the delivery apparatus. The  
 CC apparatus is used for treating occlusions or partial occlusions in  
 CC blood vessels, particularly saphenous vein grafts. The present  
 CC sequence represents a portion of an al chain of collagen, which is  
 CC given in the exemplification of the present invention.



XX Sequence 15 AA:

Query Match 100.0%; Score 80; DB 23; Length 15;  
Best Local Similarity 100.0%; Pred. No. 6.9e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPPGGIAGRGVY 15  
1 GTPPGGIAGRGVY 15

# RESULT 9

AB10111 standard; peptide: 15 AA.

AC ABB10111;

DE 12-JUL-2002 (first entry)

XX Collagen cell binding domain mimic peptide P-15.

XX Collagen; bone; repair; bone graft; tissue engineering; fibroblast;  
radiation therapy; bone damage.

OS Synthetic.

XX WO200182773-A2.

XX 08-NOV-2001.

XX 29-MAR-2001; 2001WO-US10404.

XX 28-APR-2000; 2000US-0561554.

XX (REGC ) UNIV CALIFORNIA.

XX Bhatnagar RS, Qian JY;

XX WPI; 2002-034479/04.

PT Preparation of bone repair apparatus comprises seeding at least some of  
cultured tissue cells on biologically compatible structure having  
collagen mimic and incubating seeded cells under cell growth conditions

PS Claim 7; Page 6; 23pp; English.

XX The invention relates to a bone repair apparatus that is prepared by  
growing harvested fibroblasts under cell growth conditions to form  
cultured tissue cells, seeding at least some of the cultured tissue  
cells on a biologically compatible structure having a collagen mimic, and  
incubating the seeded cells under cell growth conditions, where the  
seeded cells differentiate into an osteogenic phenotype. Methods of the  
invention are useful for preparing bone repair apparatus for use as a  
bone graft. The fibroblast cells from the recipient can be easily  
harvested with minimal invasion and trauma to the patient. By contrast to  
other methods, the fibroblast is plentiful and easily obtained with  
minimal trauma and the inventive method is able to obtain living bone  
-like cells and they, together with the biologically compatible  
structure, yield a tissue engineered bone graft. This can integrate with  
host bone when implanted in the patient, and repopulates host sites  
lacking viable cells because of disease or radiation therapy. The current  
sequence represents a collagen cell binding domain mimic peptide P-15.  
This 15 amino acid peptide has the same sequence as a particular, small  
region in the alpha(1) chain of collagen.

XX Sequence 15 AA:

Query Match 100.0%; Score 80; DB 23; Length 15;  
Best Local Similarity 100.0%; Pred. No. 6.9e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPPGGIAGRGVY 15  
1 GTPPGGIAGRGVY 15

# RESULT 10

AA92859 standard; peptide: 16 AA.

AC AAR92859;

DE 03-OCT-1996 (first entry)

XX Collagen fragment P-15 as positive control for cell adhesion.

XX Inter cellular adhesion; stimulation; inhibition; skin graft;

XX synthetic blood vessel; coating; endothelial cell; epidermal cell;

XX chemotactic attractor; wound healing; organ transplantation;

OS Synthetic.

XX Key Location/Qualifiers  
Modified-site 16 /note="C-terminal Cys residue for attaching  
peptide to a carrier protein, e.g. BSA"

XX DE4430601-A1.

XX 29-FEB-1996.

XX 22-AUG-1994; 94DE-4430601.

XX 22-AUG-1994; 94DE-4430601.

XX (BEIE ) BEIERSDORF AG.

XX Doerschner A, Eichner W, Kock K, Mielke H;

XX WPI; 1996-130242/14.

PT Peptide(s) that stimulate or inhibit cell to cell adhesion - used  
e.g. to coat synthetic blood vessels with endothelial cells, to  
prepare, or increase growth of skin grafts, to prevent thrombosis  
etc.

PS Example 1; Page 7; 18pp; German.

XX Peptides contg. the highly generic sequence AA5-AA4-AA3-AA2-AA1-(AAx)n  
where AA5 is Glu, Ser, Asp or Asn; AA4 is Leu or Ser, AA3 is Ile, Ile,  
Phe or Gly; AA2 is Asp, Leu, Asn or Ser; AA1 is Gly, Pro or Asp; AAx  
is any amino acid and n = 0 or 1 are claimed; AA5 or AA5-AA4 may be  
absent. When two or more such peptides are attached to a carrier, the  
product can be used for stimulating adhesion of eukaryotic cells in  
vitro. Particular applications include coating synthetic blood vessels  
with endothelial cells, preparing skin grafts using epithelial cells  
or stimulating wound healing. When a single peptide is used it may  
inhibit intercellular adhesion, making it useful for preventing  
thrombosis or arteriosclerosis or to suppress cancer metastases. The  
peptides can also be used as chemotactic attractors and for detecting/  
quantifying cell-cell adhesion in vitro.  
XX The present sequence is a fragment of the alpha-1 chain of collagen  
which was used as a positive control in a cell adhesion assay on the  
novel peptides.

XX Sequence 16 AA:

Query Match 100.0%; Score 80; DB 17; Length 16;  
Best Local Similarity 100.0%; Pred. No. 7.3e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;





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STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
HYPOTHETICAL: NO  
ORIGINAL SOURCE:  
ORGANISM: Homo sapiens  
TISSUE TYPE: Collagen type II  
US-08-931-820-3

Query Match 88.8%; Score 71; DB 3; Length 1060;  
Best Local Similarity 80.0%; Pred. No. 0.017;  
Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 GTPGPGIAGORGIV 15  
| | | | | : | | | | : |  
Db 788 GTPGPGIAGORGIV 802

RESULT 14  
US-08-963-825-20  
Sequence 20, Application US/08963825  
Patent No. 6110689

GENERAL INFORMATION:  
APPLICANT: Qvist, Per  
TITLE OF INVENTION: A Method for Assaying Collagen Fragments  
TITLE OF INVENTION: In Body Fluids, A Test Kit and Means for Carrying Out the  
TITLE OF INVENTION: Method and Use of the Method to Diagnose the Presence of  
NUMBER OF SEQUENCES: 21  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Darby & Darby PC  
STREET: 805 Third Avenue  
CITY: New York  
STATE: New York  
COUNTRY: USA  
ZIP: 10022

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/963,825  
FILING DATE:

CLASSIFICATION: 436  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/08/187,319  
FILING DATE: 21-JAN-1994

ATTORNEY/AGENT INFORMATION:  
NAME: Gogoris, Adda C  
REGISTRATION NUMBER: 29,714  
REFERENCE/DOCKET NUMBER: 4305/08701  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 212-527-7700  
TELEFAX: 212-753-6237  
TELEX: 236687

INFORMATION FOR SEQ ID NO: 20:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1418 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
ORIGINAL SOURCE:  
ORGANISM: Homo sapiens  
IMMEDIATE SOURCE:  
CLONE: COLLAGEN -ALPHA 1 (II)  
US-08-963-825-20

Query Match 88.8%; Score 71; DB 3; Length 1418;  
Best Local Similarity 80.0%; Pred. No. 0.023;  
Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 GTPGPGIAGORGIV 15  
| | | | | : | | | | : |  
Db 900 GTPGPGIAGORGIV 914

RESULT 15  
US-09-010-999-1  
Sequence 1, Application US/09010999  
Patent No. 6132976

GENERAL INFORMATION:  
APPLICANT: Poole, Anthony R.  
APPLICANT: Hollander, Anthony P.  
APPLICANT: Billingham, R. C.  
TITLE OF INVENTION: IMMUNOASSAYS FOR THE MEASUREMENT OF  
TITLE OF INVENTION: COLLAGEN DENATURATION AND CLEAVAGE IN CARTILAGE  
NUMBER OF SEQUENCES: 16  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Foley & Lardner  
STREET: 3000 K Street, N.W., Suite 500  
CITY: Washington  
STATE: D.C.  
COUNTRY: USA  
ZIP: 20007-5109

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/010,999  
FILING DATE: 22-JAN-1998  
CLASSIFICATION: 4335

PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 08/448,501  
FILING DATE: 17-JUL-1995  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 07/984,123  
FILING DATE: 04-DEC-1992

ATTORNEY/AGENT INFORMATION:  
NAME: Bent, Stephen A.  
REGISTRATION NUMBER: 29,768  
REFERENCE/DOCKET NUMBER: 032931/0212  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (202)672-5300  
TELEFAX: (202)672-5399  
TELEX: 904136

INFORMATION FOR SEQ ID NO: 1:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1418 amino acids  
TYPE: amino acid  
STRANDEDNESS:  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
ORIGINAL SOURCE:  
ORGANISM: Human Type II Collagen  
US-09-010-999-1

Query Match 88.8%; Score 71; DB 3; Length 1418;  
Best Local Similarity 80.0%; Pred. No. 0.023;  
Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;  
OY 1 GTPGPGIAGORGIV 15  
| | | | | : | | | | : |  
Db 900 GTPGPGIAGORGIV 914

Search completed: August 29, 2003, 18:30:11  
Job time: 30 secs

;; TITLE OF INVENTION: Disorders Associated with the Metabolism of  
;; NUMBER OF SEQUENCES: 21  
;; CORRESPONDENCE ADDRESS:  
;; ADDRESSEE: Darby & Darby PC  
;; STREET: 805 Third Avenue  
;; CITY: New York  
;; STATE: New York  
;; COUNTRY: USA  
;; ZIP: 10022  
;; COMPUTER READABLE FORM:  
;; MEDIUM TYPE: Floppy disk  
;; COMPUTER: IBM PC compatible  
;; OPERATING SYSTEM: PC-DOS/MS-DOS  
;; SOFTWARE: Patent Release #1.0, Version #1.25  
;; CURRENT APPLICATION DATA:  
;; APPLICATION NUMBER: US/09/548,608  
;; FILING DATE:  
;; CLASSIFICATION:  
;; PRIOR APPLICATION DATA:  
;; APPLICATION NUMBER: 08/187,319  
;; FILING DATE:  
;; ATTORNEY/AGENT INFORMATION:  
;; NAME: Gogoris, Adda C  
;; REGISTRATION NUMBER: 29,714  
;; REFERENCE/DOCKET NUMBER: 4305/08701  
;; TELECOMMUNICATION INFORMATION:  
;; TELEPHONE: 212-527-7700  
;; TELEFAX: 212-753-6237  
;; TELEX: 236687  
;; INFORMATION FOR SEQ ID NO: 18:  
;; SEQUENCE CHARACTERISTICS:  
;; LENGTH: 1341 amino acids  
;; TYPE: amino acid  
;; TOPOLOGY: linear  
;; MOLECULE TYPE: protein  
;; ORIGINAL SOURCE:  
;; ORGANISM: Homo sapiens  
;; IMMEDIATE SOURCE:  
;; CLONE: COLLAGEN ALPHA 1 (I)  
;; US-09-548-608-18

Query Match 100.0%; Score 80; DB 4; Length 1341;  
Best Local Similarity 100.0%; Pred. No. 0.0011;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGV 15  
DB 823 GTPGPGIAGRGV 837

RESULT 11  
US-09-585-887-9  
; Sequence 9, Application US/09585887  
; Patent No. 6413742  
; GENERAL INFORMATION:  
; APPLICANT: Olsen, David R  
; APPLICANT: Chang, Robert  
; APPLICANT: McMullin, Hugh  
; APPLICANT: Hitzeman, Ronald A.  
; APPLICANT: Chisholm, George  
; TITLE OF INVENTION: NOVEL METHODS FOR THE PRODUCTION OF GELATIN AND  
; TITLE OF INVENTION: FULL-LENGTH TRIPLE HELICAL COLLAGEN IN RECOMBINANT  
; TITLE OF INVENTION: CELLS  
; FILE REFERENCE: 225002030400  
; CURRENT APPLICATION NUMBER: US/09/585,887  
; CURRENT FILING DATE: 2000-05-31  
; PRIOR APPLICATION NUMBER: 09/289,578  
; PRIOR FILING DATE: 1999-04-09  
; PRIOR APPLICATION NUMBER: 60/084,828  
; PRIOR FILING DATE: 1998-05-08  
; NUMBER OF SEQ ID NOS: 11  
; SOFTWARE: Patentin Ver. 2.0  
; SEQ ID NO 9

;; LENGTH: 1461  
;; TYPE: PRT  
;; ORGANISM: Homo sapiens  
;; US-09-585-887-9

Query Match 100.0%; Score 80; DB 4; Length 1461;  
Best Local Similarity 100.0%; Pred. No. 0.0012;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGV 15  
DB 944 GTPGPGIAGRGV 958

RESULT 12  
US-09-289-578-9  
; Sequence 9, Application US/09289578  
; Patent No. 6428978  
; GENERAL INFORMATION:  
; APPLICANT: Olsen, David R  
; APPLICANT: Chang, Robert  
; APPLICANT: McMullin, Hugh  
; APPLICANT: Hitzeman, Ronald A.  
; APPLICANT: Chisholm, George  
; TITLE OF INVENTION: NOVEL METHODS FOR THE PRODUCTION OF GELATIN AND  
; TITLE OF INVENTION: FULL-LENGTH TRIPLE HELICAL COLLAGEN IN RECOMBINANT  
; TITLE OF INVENTION: CELLS  
; FILE REFERENCE: 225002030400  
; CURRENT APPLICATION NUMBER: US/09/289,578  
; CURRENT FILING DATE: 1999-04-10  
; PRIOR APPLICATION NUMBER: 60/084,828  
; PRIOR FILING DATE: 1998-05-08  
; NUMBER OF SEQ ID NOS: 11  
; SOFTWARE: Patentin Ver. 2.0  
; SEQ ID NO 9  
; LENGTH: 1461  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
; US-09-289-578-9

Query Match 100.0%; Score 80; DB 4; Length 1461;  
Best Local Similarity 100.0%; Pred. No. 0.0012;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGV 15  
DB 944 GTPGPGIAGRGV 958

RESULT 13  
US-08-931-820-3  
; Sequence 3, Application US/08931820  
; Patent No. 6010863  
; GENERAL INFORMATION:  
; APPLICANT:  
; TITLE OF INVENTION: Assay for collagen degradation  
; NUMBER OF SEQUENCES: 4  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/931,820  
; FILING DATE:  
; CLASSIFICATION: 435  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: EP 96202596.1  
; FILING DATE:  
; INFORMATION FOR SEQ ID NO: 3:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 1060 amino acids  
; TYPE: amino acid

MOLECULE TYPE: protein  
ORIGINAL SOURCE:  
ORGANISM: Homo sapiens  
IMMEDIATE SOURCE:  
CLONE: COLLAGEN ALPHA 1 (1)  
US-08-963-825-18

Query Match 100.0%; Score 80; DB 3; Length 1341;  
Best Local Similarity 100.0%; Pred. No. 0.0011;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGORGVV 15  
|||||  
Db 823 GTPGPGIAGORGVV 837

RESULT 8  
US-09-500-811-18  
Sequence 18, Application US/09500811  
Patent No. 6323314  
GENERAL INFORMATION:  
APPLICANT: Ovisst, Per  
APPLICANT: Bonde, Martin  
TITLE OF INVENTION: A Method for Assaying Collagen Fragments  
TITLE OF INVENTION: In Body Fluids, A Test Kit and Means for Carrying Out the  
TITLE OF INVENTION: Method and Use of the Method to Diagnose the Presence of  
TITLE OF INVENTION: Disorders Associated with the Metabolism of  
NUMBER OF SEQUENCES: 21  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Darby & Darby PC  
STREET: 805 Third Avenue  
CITY: New York  
STATE: New York  
COUNTRY: USA  
ZIP: 10022  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/500,811  
TELEPHONE: 212-527-7700  
TELEFAX: 212-753-6237  
FILING DATE:  
CLASSIFICATION:  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/187,319  
FILING DATE:  
ATTORNEY/AGENT INFORMATION:  
NAME: Gogoris, Adda C  
REGISTRATION NUMBER: 29,714  
REFERENCE/DOCKET NUMBER: 4305/08701  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 212-527-7700  
TELEFAX: 212-753-6237  
INFORMATION FOR SEQ ID NO: 18:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1341 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
ORIGINAL SOURCE:  
ORGANISM: Homo sapiens  
IMMEDIATE SOURCE:  
CLONE: COLLAGEN ALPHA 1 (1)  
US-09-500-811-18

Query Match 100.0%; Score 80; DB 4; Length 1341;  
Best Local Similarity 100.0%; Pred. No. 0.0011;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGORGVV 15  
|||||

Db 823 GTPGPGIAGORGVV 837

RESULT 9  
US-09-570-573-18  
Sequence 18, Application US/09570573  
Patent No. 6342361  
GENERAL INFORMATION:  
APPLICANT: Ovisst, Per  
APPLICANT: Bonde, Martin  
TITLE OF INVENTION: A Method for Assaying Collagen Fragments  
TITLE OF INVENTION: In Body Fluids, A Test Kit and Means for Carrying Out the  
TITLE OF INVENTION: Method and Use of the Method to Diagnose the Presence of  
TITLE OF INVENTION: Disorders Associated with the Metabolism of  
NUMBER OF SEQUENCES: 21  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Darby & Darby PC  
STREET: 805 Third Avenue  
CITY: New York  
STATE: New York  
COUNTRY: USA  
ZIP: 10022  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/570,573  
FILING DATE:  
CLASSIFICATION:  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 08/187,319  
FILING DATE:  
ATTORNEY/AGENT INFORMATION:  
NAME: Gogoris, Adda C  
REGISTRATION NUMBER: 29,714  
REFERENCE/DOCKET NUMBER: 4305/08701  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 212-527-7700  
TELEFAX: 212-753-6237  
INFORMATION FOR SEQ ID NO: 18:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1341 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
ORIGINAL SOURCE:  
ORGANISM: Homo sapiens  
IMMEDIATE SOURCE:  
CLONE: COLLAGEN ALPHA 1 (1)  
US-09-570-573-18

Query Match 100.0%; Score 80; DB 4; Length 1341;  
Best Local Similarity 100.0%; Pred. No. 0.0011;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGORGVV 15  
|||||

Db 823 GTPGPGIAGORGVV 837

RESULT 10  
US-09-548-608-18  
Sequence 18, Application US/09548608  
Patent No. 6355442  
GENERAL INFORMATION:  
APPLICANT: Ovisst, Per  
APPLICANT: Bonde, Martin  
TITLE OF INVENTION: A Method for Assaying Collagen Fragments  
TITLE OF INVENTION: In Body Fluids, A Test Kit and Means for Carrying Out the  
TITLE OF INVENTION: Method and Use of the Method to Diagnose the Presence of

EARLIER FILING DATE: 1998-09-10  
NUMBER OF SEQ ID NOS: 23  
SOFTWARE: PatentIn Ver. 2.1  
SEQ ID NO: 3  
LENGTH: 25  
TYPE: PRT  
ORGANISM: Homo sapiens  
FEATURE:  
OTHER INFORMATION: COL1A1 Binding Peptides  
US-09-517-866-3

Query Match 100.0%; Score 80; DB 4; Length 25;  
Best Local Similarity 100.0%; Pred. No. 2.3e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCGV 15  
|||||  
DB 9 GTPGPGIAGGRCGV 23

RESULT 5  
US-09-219-849-49  
Sequence 49, Application US/09219849  
Patent No. 6150081  
GENERAL INFORMATION:  
APPLICANT: VAN HEERDE, GEORGE V.  
APPLICANT: VAN RIJN, ALEXIS C.  
APPLICANT: BOUWSTRA, JAN B.  
APPLICANT: DE WOLF, FREDERIK A.  
APPLICANT: MOORROEK, ANDREAS  
APPLICANT: WERTEN, MARC W.T.  
APPLICANT: WIND, RICHELIE D.  
APPLICANT: VAN DEN BOSCH, TANJA J.  
TITLE OF INVENTION: SILVER HALIDE EMULSIONS WITH RECOMBINANT COLLAGEN  
TITLE OF INVENTION: SUITABLE FOR PHOTOGRAPHIC APPLICATION AND ALSO THE  
FILE REFERENCE: 2728-2  
CURRENT APPLICATION NUMBER: US/09/219,849  
CURRENT FILING DATE: 1998-12-23  
NUMBER OF SEQ ID NOS: 50  
SOFTWARE: PatentIn Ver. 2.1  
SEQ ID NO 49  
LENGTH: 822  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: Description of Artificial Sequence: Synthetic  
US-09-219-849-49

Query Match 100.0%; Score 80; DB 3; Length 822;  
Best Local Similarity 100.0%; Pred. No. 0.00069;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCGV 15  
|||||  
DB 768 GTPGPGIAGGRCGV 782

RESULT 6  
US-08-931-820-1  
Sequence 1, Application US/08931820  
Patent No. 6010863  
GENERAL INFORMATION:  
APPLICANT:  
TITLE OF INVENTION: Assay for collagen degradation  
NUMBER OF SEQUENCES: 4  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)  
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/931,820  
FILING DATE:  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: EP 96202596.1  
FILING DATE:  
INFORMATION FOR SEQ ID NO: 1:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1057 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
HYPOTHEICAL: NO  
ANTI-SENSE: NO  
ORIGINAL SOURCE:  
ORGANISM: Homo sapiens  
TISSUE TYPE: Collagen type I  
US-08-931-820-1

Query Match 100.0%; Score 80; DB 3; Length 1057;  
Best Local Similarity 100.0%; Pred. No. 0.00088;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCGV 15  
|||||  
DB 786 GTPGPGIAGGRCGV 800

RESULT 7  
US-08-963-825-18  
Sequence 18, Application US/08963825  
Patent No. 6110689  
GENERAL INFORMATION:  
APPLICANT: Qvist, Per  
APPLICANT: Bonde, Martin  
TITLE OF INVENTION: A Method for Assaying Collagen Fragments  
TITLE OF INVENTION: In Body Fluids, A Test Kit and Means for Carrying Out the  
TITLE OF INVENTION: Method and Use of the Method to Diagnose the Presence of  
TITLE OF INVENTION: Disorders Associated with the Metabolism of  
NUMBER OF SEQUENCES: 21  
CORRESPONDENCE ADDRES:  
ADDRESSEE: Darby & Darby PC  
STREET: 805 Third Avenue  
CITY: New York  
STATE: New York  
COUNTRY: USA  
ZIP: 10022  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/963,825  
FILING DATE:  
CLASSIFICATION: 436  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/08/187,319  
FILING DATE: 21-JAN-1994  
ATTORNEY/AGENT INFORMATION:  
NAME: Gogoris, Adia C  
REGISTRATION NUMBER: 29,714  
REFERENCE/DOCKET NUMBER: 4305/08701  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 212-527-7700  
TELEFAX: 212-753-6237  
TELEX: 236687  
INFORMATION FOR SEQ ID NO: 18:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1341 amino acids  
TYPE: amino acid  
TOPOLOGY: linear



Query Match 100.0%; Score 80; DB 2; Length 15;  
Best Local Similarity 100.0%; Pred. No. 1.4e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 GTGPGGIAGRGV 15  
Db 1 GTGPGGIAGRGV 15

## RESULT 2

US-09-328-347A-1  
; Sequence 1, Application US/09328347A  
; Patent No. 6268348  
; GENERAL INFORMATION:  
; APPLICANT: Bhatnagar, Rajendra S.  
; TITLE OF INVENTION: SYNTHETIC COMPOUNDS AND COMPOSITIONS  
; TITLE OF INVENTION: WITH ENHANCED CELL BINDING  
; NUMBER OF SEQUENCES: 14  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Majestic, Parsons, Siebert & Hsue P.C.  
; STREET: Four Embarcadero Center, Suite 1100  
; CITY: San Francisco  
; STATE: California  
; COUNTRY: U.S.A.  
; ZIP: 94111-4106  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patentin Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/09/328,347A  
; FILING DATE: 08-JUN-1999  
; CLASSIFICATION: 435  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 08/859,610  
; FILING DATE: 20-MAY-1997  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 08/278,878  
; FILING DATE: 22-JUL-1994  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 07/804,782  
; FILING DATE: 09-DEC-1991  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 07/393,621  
; FILING DATE: 14-AUG-1989  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Siebert, J. Suzanne  
; REGISTRATION NUMBER: 28,758  
; REFERENCE/DOCKET NUMBER: 2500,066US5  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 415-248-5500  
; TELEFAX: 415-362-5418.  
; INFORMATION FOR SEQ ID NO: 1:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 15 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; HYPOTHEICAL: NO  
; ANTI-SENSE: NO  
; US-09-328-347A-1

Query Match 100.0%; Score 80; DB 3; Length 15;  
Best Local Similarity 100.0%; Pred. No. 1.4e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 GTGPGGIAGRGV 15  
Db 1 GTGPGGIAGRGV 15

## RESULT 3

US-09-010-999-9  
; Sequence 9, Application US/09010999  
; Patent No. 6132976  
; GENERAL INFORMATION:  
; APPLICANT: Poole, Anthony R.  
; APPLICANT: Hollander, Anthony P.  
; APPLICANT: Billingham, R. C.  
; TITLE OF INVENTION: IMMUNOASSAYS FOR THE MEASUREMENT OF  
; TITLE OF INVENTION: COLLAGEN DENATURATION AND CLEAVAGE IN CARTILAGE  
; NUMBER OF SEQUENCES: 16  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Foley & Lardner  
; STREET: 3000 K Street, N.W., Suite 500  
; CITY: Washington  
; STATE: D.C.  
; COUNTRY: USA  
; ZIP: 20007-5109  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patentin Release #1.0, Version #1.30  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/09/010,999  
; FILING DATE: 22-JAN-1998  
; CLASSIFICATION: 4335  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 08/448,501  
; FILING DATE: 17-JUL-1995  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 07/984,123  
; FILING DATE: 04-DEC-1992  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Bent, Stephen A.  
; REGISTRATION NUMBER: 29,768  
; REFERENCE/DOCKET NUMBER: 032931/0212  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (202)672-5300  
; TELEFAX: (202)672-5399  
; TELERX: 904136  
; INFORMATION FOR SEQ ID NO: 9:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 19 amino acids  
; TYPE: amino acid  
; STRANDEDNESS:  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; US-09-010-999-9

Query Match 100.0%; Score 80; DB 3; Length 19;  
Best Local Similarity 100.0%; Pred. No. 1.7e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 GTGPGGIAGRGV 15  
Db 4 GTGPGGIAGRGV 18

## RESULT 4

US-09-517-866-3  
; Sequence 3, Application US/09517866  
; Patent No. 6472504  
; GENERAL INFORMATION:  
; APPLICANT: PROCKOP, Darwin J.  
; APPLICANT: FERTALA, Andrez J.  
; TITLE OF INVENTION: INHIBITORS OF COLLAGEN ASSEMBLY  
; FILE REFERENCE: 209598.0111/2801  
; CURRENT APPLICATION NUMBER: US/09/517,866  
; EARLIER FILING DATE: 2000-03-03  
; EARLIER APPLICATION NUMBER: 60/058,353  
; EARLIER FILING DATE: 1997-09-10  
; EARLIER APPLICATION NUMBER: PCT/US98/18838

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: August 29, 2003, 18:25:20 ; Search time 29 seconds  
(without alignments)  
21.885 Million cell updates/sec

Title: US-09-935-417-1  
Perfect score: 80  
Sequence: 1 GTPGPGIAGQRCVV 15

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 328717 seqs, 42310858 residues

Total number of hits satisfying chosen parameters: 328717

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

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3: /cgn2\_6/ptodata/1/1aa/6A.COMB.pep:\*  
4: /cgn2\_6/ptodata/1/1aa/6B.COMB.pep:\*  
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6: /cgn2\_6/ptodata/1/1aa/Backfiles1.pep:\*

Pred. No. is the number of results predicted by chance to have a  
Perfect greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	80	100.0	15	2	US-08-859-610A-1
2	80	100.0	15	3	US-09-328-347A-1
3	80	100.0	19	3	US-09-010-999-9
4	80	100.0	25	4	US-09-517-866-3
5	80	100.0	822	3	US-09-219-849-49
6	80	100.0	1057	3	US-08-931-820-1
7	80	100.0	1341	3	US-08-963-825-18
8	80	100.0	1341	4	US-09-500-811-18
9	80	100.0	1341	4	US-09-570-573-18
10	80	100.0	1341	4	US-09-548-608-18
11	80	100.0	1461	4	US-09-585-887-9
12	80	100.0	1461	4	US-09-289-578-9
13	80	100.0	1060	3	US-08-931-820-3
14	80	100.0	1418	3	US-08-963-825-20
15	80	100.0	1418	3	US-09-010-999-1
16	80	100.0	1418	4	US-09-500-811-20
17	80	100.0	1418	4	US-09-570-573-20
18	80	100.0	1418	4	US-09-548-608-20
19	80	100.0	1442	5	PCT-US95-02251-12
20	80	100.0	1442	5	PCT-US95-02251-12
21	62	77.5	19	3	US-09-010-999-8
22	61	76.2	12	1	US-08-330-599-16
23	57	71.2	1057	3	US-08-931-820-4
24	57	71.2	1078	3	US-08-963-825-21
25	57	71.2	1078	4	US-09-500-811-21
26	57	71.2	1078	4	US-09-570-573-21
27	57	71.2	1078	4	US-09-548-608-21

28	56	70.0	19	3	US-09-010-999-10	Sequence 10, Appl
29	56	70.0	1024	3	US-08-931-820-2	Sequence 2, Appl
30	56	70.0	1366	3	US-08-963-825-19	Sequence 19, Appl
31	56	70.0	1366	4	US-09-500-811-19	Sequence 19, Appl
32	56	70.0	1366	4	US-09-570-573-19	Sequence 19, Appl
33	56	70.0	1366	4	US-09-548-608-19	Sequence 19, Appl
34	56	70.0	1366	4	US-09-585-887-10	Sequence 10, Appl
35	56	70.0	1366	4	US-09-289-578-10	Sequence 10, Appl
36	54	67.5	228	3	US-09-219-849-38	Sequence 38, Appl
37	54	67.5	279	3	US-09-010-999-2	Sequence 2, Appl
38	54	67.5	417	1	US-08-175-155-69	Sequence 69, Appl
39	54	67.5	417	1	US-08-477-509B-104	Sequence 104, App
40	54	67.5	417	1	US-08-642-255-102	Sequence 102, App
41	54	67.5	417	2	US-08-707-237A-76	Sequence 76, Appl
42	54	67.5	417	3	US-08-482-085B-104	Sequence 104, App
43	54	67.5	417	4	US-09-444-791A-104	Sequence 104, App
44	54	67.5	829	1	US-08-642-255-132	Sequence 132, App
45	54	67.5	829	1	US-08-397-633A-53	Sequence 53, Appl

#### ALIGNMENTS

RESULT 1  
US-08-859-610A-1  
; Sequence 1, Application US/08859610A  
; Patent No. 5958428  
; GENERAL INFORMATION:  
; APPLICANT: Bhatnagar, Rajendra S.  
; TITLE OF INVENTION: SYNTHETIC COMPOUNDS AND COMPOSITIONS  
; TITLE OF INVENTION: WITH ENHANCED CELL BINDING  
; NUMBER OF SEQUENCES: 14  
; CORRESPONDENCE ADDRESS:  
; ADDRESS: Majestic, Parsons, Siebert & Hae P.C.  
; STREET: Four Embarcadero Center, Suite 1100  
; CITY: San Francisco  
; STATE: California  
; COUNTRY: U.S.A.  
; ZIP: 94111-4106  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patentin Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/859,610A  
; FILING DATE: 20-MAY-1997  
; CLASSIFICATION: 435  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 08/278,878  
; FILING DATE: 22-JUL-1994  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 07/804,782  
; FILING DATE: 09-DEC-1991  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 07/393,621  
; FILING DATE: 14-AUG-1989  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Siebert, J. Suzanne  
; REGISTRATION NUMBER: 28,758  
; REFERENCE/DOCKET NUMBER: 2500.066US4  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 415-248-5500  
; TELEFAX: 415-362-5418  
; INFORMATION FOR SEQ ID NO: 1:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 15 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; HYPOTHETICAL: NO  
; ANTI-SENSE: NO  
; US-08-859-610A-1

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OM protein - protein search, using sw model

Run on: August 29, 2003, 18:26:40 ; Search time 25 Seconds

(without alignments)  
82.069 Million cell updates/sec

Title: US-09-935-417-1

Sequence: 1 GTPGPGIAGRGV 15

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Gapop 10.0, Gapext 0.5

Searched: 510680 seqs, 136781880 residues

Total number of hits satisfying chosen parameters: 510680

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Published Applications AA:\*

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5: /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	80	100.0	15	9	US-09-113-696B-17
2	80	100.0	15	9	US-09-816-737-1
3	80	100.0	15	9	US-09-935-417-1
4	80	100.0	15	15	US-10-133-289-1
5	80	100.0	15	15	US-10-176-401-1
6	80	100.0	15	15	US-10-017-193-1
7	80	100.0	25	15	US-10-279-991-3
8	80	100.0	1341	15	US-10-058-124-18
9	80	100.0	1464	12	US-10-301-822-38
10	80	100.0	1464	15	US-10-060-036-159
11	80	100.0	1464	15	US-10-171-311-36
12	80	100.0	1464	15	US-10-216-705-21
13	80	100.0	1464	15	US-10-149-352-2
14	80	100.0	1464	15	US-10-177-293-65
15	71	88.8	234	11	US-09-895-674-1

16	71	88.8	714	15	US-10-233-885-44	Sequence 44, Appl
17	71	88.8	714	15	US-10-231-581-44	Sequence 44, Appl
18	71	88.8	1014	12	US-10-194-441A-1	Sequence 1, Appl
19	71	88.8	1014	12	US-10-194-441A-48	Sequence 48, Appl
20	71	88.8	1418	15	US-10-058-124-20	Sequence 20, Appl
21	60	75.0	459	9	US-09-789-561-97	Sequence 97, Appl
22	58	72.5	1496	12	US-10-301-822-35	Sequence 35, Appl
23	58	72.5	1496	15	US-10-177-293-70	Sequence 70, Appl
24	57	71.2	1078	15	US-10-058-124-21	Sequence 21, Appl
25	57	71.2	1466	12	US-10-301-822-33	Sequence 33, Appl
26	57	71.2	1466	15	US-10-177-293-68	Sequence 68, Appl
27	56	70.0	674	9	US-09-925-299-979	Sequence 979, Appl
28	56	70.0	674	11	US-09-925-299-979	Sequence 979, Appl
29	56	70.0	1366	12	US-10-301-822-31	Sequence 31, Appl
30	56	70.0	1366	15	US-10-171-311-38	Sequence 38, Appl
31	56	70.0	1366	15	US-10-058-124-19	Sequence 19, Appl
32	54	67.5	33	9	US-09-864-761-39163	Sequence 39163, A
33	54	67.5	417	15	US-10-096-986-104	Sequence 104, App
34	54	67.5	837	15	US-10-096-986-103	Sequence 103, App
35	53	66.2	61	15	US-10-096-986-102	Sequence 102, App
36	53	66.2	680	15	US-10-177-293-59	Sequence 59, Appl
37	53	66.2	742	15	US-10-203-860-4	Sequence 4, Appl
38	52	65.0	403	9	US-09-925-302-689	Sequence 689, App
39	52	65.0	520	10	US-09-978-295A-614	Sequence 614, App
40	52	65.0	520	10	US-09-978-697-614	Sequence 614, App
41	52	65.0	520	10	US-09-978-122A-614	Sequence 614, App
42	52	65.0	520	10	US-09-999-832A-614	Sequence 614, App
43	52	65.0	520	11	US-09-978-189-614	Sequence 614, App
44	52	65.0	520	11	US-09-978-608A-614	Sequence 614, App
45	52	65.0	520	11	US-09-978-585A-614	Sequence 614, App

## ALIGNMENTS

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RESULT 1
US-09-113-696B-17
; Sequence 17, Application US/09113696B
; Patent No. US20020010134A1
; GENERAL INFORMATION:
; APPLICANT: Bhatnagar, Rajendra S.
; APPLICANT: Qian, Jing Jing
; TITLE OF INVENTION: PEPTIDE COMPOSITIONS MIMICKING TGF-BETA
; FILE REFERENCE: 6510-215CIP2
; CURRENT APPLICATION NUMBER: US/09/113,696B
; CURRENT FILING DATE: 1998-07-10
; PRIOR APPLICATION NUMBER: 08/742,256
; PRIOR FILING DATE: 1996-10-31
; PRIOR APPLICATION NUMBER: 08/431,954
; PRIOR FILING DATE: 1995-05-01
; NUMBER OF SEQ ID NOS: 42
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 17
; LENGTH: 15
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: TGF-beta mimic
US-09-113-696B-17
Query Match 100.0%, Score 80; DB 9; Length 15;
Best Local Similarity 100.0%; Pred. No. 3.8e-05;
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 GTPGPGIAGRGV 15
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Db 1 GTPGPGIAGRGV 15

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RESULT 2  
US-09-816-737-1

Sequence 1, Application US/09616737  
Patent No. US20020037853A1  
GENERAL INFORMATION:  
APPLICANT: Bhatnagar, Rajendra S.  
TITLE OF INVENTION: "Synthetic Compounds and Compositions  
FILE REFERENCE: 06510223CON2  
CURRENT APPLICATION NUMBER: US/09/816,737  
CURRENT FILING DATE: 2001-03-23  
PRIOR APPLICATION NUMBER: 09/328,347  
PRIOR FILING DATE: 1999-06-08  
PRIOR APPLICATION NUMBER: 08/859,610  
PRIOR FILING DATE: 1997-05-20  
PRIOR APPLICATION NUMBER: 08/278,878  
PRIOR FILING DATE: 1994-07-22  
PRIOR APPLICATION NUMBER: 07/804,782  
PRIOR FILING DATE: 1991-12-09  
PRIOR APPLICATION NUMBER: 07/393,621  
PRIOR FILING DATE: 1989-08-14  
NUMBER OF SEQ ID NOS: 14  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 1  
LENGTH: 15  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: synthetic peptide  
US-09-816-737-1

Query Match 100.0%; Score 80; DB 9; Length 15;  
Best Local Similarity 100.0%; Pred. No. 3.8e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 GTPGPGIAGGQGVV 15  
|||||

DB 1 GTPGPGIAGGQGVV 15

RESULT 3  
US-09-935-417-1  
Sequence 1, Application US/09935417  
Patent No. US20020062145A1  
GENERAL INFORMATION:  
APPLICANT: Rudakov, Leon V.  
APPLICANT: Imran, Mtr A.  
APPLICANT: Dinh, Linh  
APPLICANT: Davidian, Ara  
APPLICANT: Larkin, Kevin  
TITLE OF INVENTION: Composite Expandable Device with Polymeric Covering and Bioactive  
FILE REFERENCE: 52200-8006,US01  
CURRENT APPLICATION NUMBER: US/09/935,417  
CURRENT FILING DATE: 2001-08-22  
PRIOR APPLICATION NUMBER: US 09/385,691  
PRIOR FILING DATE: 1999-08-30  
NUMBER OF SEQ ID NOS: 1  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 1  
LENGTH: 15  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: portion of a1 chain of collagen  
US-09-935-417-1

Query Match 100.0%; Score 80; DB 9; Length 15;  
Best Local Similarity 100.0%; Pred. No. 3.8e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 GTPGPGIAGGQGVV 15  
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DB 1 GTPGPGIAGGQGVV 15

RESULT 4  
US-10-133-289-1  
Sequence 1, Application US/10133289  
Publication No. US20030077825A1  
GENERAL INFORMATION:  
APPLICANT: Bhatnagar, Rajendra S.  
APPLICANT: Jing, Jing Qian  
TITLE OF INVENTION: Structures Useful for Bone Engineering  
FILE REFERENCE: UCAL224  
CURRENT APPLICATION NUMBER: US/10/133,289  
CURRENT FILING DATE: 2002-04-25  
PRIOR APPLICATION NUMBER: US/09/561,554  
PRIOR FILING DATE: 2000-04-28  
NUMBER OF SEQ ID NOS: 2  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 1  
LENGTH: 15  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: Synthetic Peptide  
US-10-133-289-1

Query Match 100.0%; Score 80; DB 15; Length 15;  
Best Local Similarity 100.0%; Pred. No. 3.8e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 GTPGPGIAGGQGVV 15  
|||||

DB 1 GTPGPGIAGGQGVV 15

RESULT 5  
US-10-176-401-1  
Sequence 1, Application US/10176401  
Publication No. US20030103960A1  
GENERAL INFORMATION:  
APPLICANT: Philippart, Pierre  
APPLICANT: Brasseur, Michele  
TITLE OF INVENTION: Sealant and bone generating product  
FILE REFERENCE: 402119  
CURRENT APPLICATION NUMBER: US/10/176,401  
CURRENT FILING DATE: 2002-06-21  
NUMBER OF SEQ ID NOS: 1  
SOFTWARE: PatentIn version 3.1  
SEQ ID NO 1  
LENGTH: 15  
TYPE: PRT  
ORGANISM: Homo sapiens  
FEATURE:  
NAME/KEY: PEPTIDE  
LOCATION: (1)..(15)  
OTHER INFORMATION: residue 766 to 780 from human collagen type 1  
PUBLICATION INFORMATION:  
AUTHORS: Bhatnagar RS, Qian JY, Gough CA  
TITLE: The role in cell binding of a beta-bend within the triple helical region  
TITLE: in collagen alpha 1 (I) chain: structural and biological evidence for  
TITLE: conformational tautomerism on fiber surface  
JOURNAL: Journal of Biomolecular structure and dynamics  
VOLUME: 14  
ISSUE: 5  
PAGES: 547-560  
DATE: 1997-04-01  
DATABASE ENTRY DATE:  
RELEVANT RESIDUES: (766)..(780)  
US-10-176-401-1

Query Match 100.0%; Score 80; DB 15; Length 15;  
Best Local Similarity 100.0%; Pred. No. 3.8e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCGV 15  
|||||  
Db 1 GTPGPGIAGGRCGV 15

## RESULT 6

US-10-017-193-1  
; Sequence 1, Application US/10017193  
; Publication No. US20030113478A1  
; GENERAL INFORMATION:  
; APPLICANT: Dang, Mai Huong  
; APPLICANT: Chiu, Phillip  
; TITLE OF INVENTION: Surface Coating Method and Coated Device  
; FILE REFERENCE: 52200-8010  
; CURRENT APPLICATION NUMBER: US/10/017,193  
; CURRENT FILING DATE: 2001-12-12  
; NUMBER OF SEQ ID NOS: 10  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO: 1  
; LENGTH: 15  
; TYPE: PRT  
; ORGANISM: Artificial Sequence  
; FEATURE:  
; OTHER INFORMATION: attachment peptide from collagen  
US-10-017-193-1

Query Match 100.0%; Score 80; DB 15; Length 15;  
Best Local Similarity 100.0%; Pred. No. 3.8e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCGV 15  
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Db 1 GTPGPGIAGGRCGV 15

## RESULT 7

US-10-279-991-3  
; Sequence 3, Application US/10279991  
; Publication No. US20030087315A1  
; GENERAL INFORMATION:  
; APPLICANT: PROCKOP, DARWIN J.  
; APPLICANT: PERTALIA, ANDREJ  
; TITLE OF INVENTION: INHIBITORS OF COLLAGEN ASSEMBLY  
; FILE REFERENCE: 053844-5001-01  
; CURRENT APPLICATION NUMBER: US/10/279,991  
; CURRENT FILING DATE: 2002-10-24  
; PRIOR APPLICATION NUMBER: 09/517,866  
; PRIOR FILING DATE: 2000-03-03  
; PRIOR APPLICATION NUMBER: 60/058,353  
; PRIOR FILING DATE: 1997-09-10  
; PRIOR APPLICATION NUMBER: PCT/US98/18838  
; PRIOR FILING DATE: 1998-09-10  
; NUMBER OF SEQ ID NOS: 23  
; SOFTWARE: PatentIn version 3.1  
; SEQ ID NO: 3  
; LENGTH: 25  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-10-279-991-3

Query Match 100.0%; Score 80; DB 15; Length 25;  
Best Local Similarity 100.0%; Pred. No. 6.3e-05;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCGV 15  
|||||  
Db 9 GTPGPGIAGGRCGV 23

RESULT 8  
US-10-058-124-18  
; Sequence 18, Application US/10058124  
; Publication No. US20030119058A1

## GENERAL INFORMATION:

APPLICANT: Qvist, Per

Bonde, Martin

TITLE OF INVENTION: A Method for Assaying Collagen Fragments  
in Body Fluids, A Test Kit and Means for Carrying Out th  
Method and Use of the Method to Diagnose the Presence of  
Disorders Associated with the Metabolism of

NUMBER OF SEQUENCES: 21

CORRESPONDENCE ADDRESS:

ADDRESSEE: Dady & Dady PC

STREET: 805 Third Avenue

CITY: New York

STATE: New York

COUNTRY: USA

ZIP: 10022

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/058,124

FILING DATE: 29-Jan-2002

CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/570,573

FILING DATE: 2002-MAY-12

APPLICATION NUMBER: 08/187,319

FILING DATE: <Unknown>

ATTORNEY/AGENT INFORMATION:

NAME: Gogosis, Adda C

REGISTRATION NUMBER: 29,714

REFERENCE/DOCKET NUMBER: 4305/08701

TELECOMMUNICATION INFORMATION:

TELEPHONE: 212-527-7700

TELEFAX: 212-753-6237

TELEX: 236687

INFORMATION FOR SEQ ID NO: 18:

SEQUENCE CHARACTERISTICS:

LENGTH: 1341 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

ORIGINAL SOURCE:

ORGANISM: Homo sapiens

IMMEDIATE SOURCE:

CLONE: COLLAGEN ALPHA 1 (I)

SEQUENCE DESCRIPTION: SEQ ID NO: 18:

US-10-058-124-18

Query Match 100.0%; Score 80; DB 15; Length 1341;  
Best Local Similarity 100.0%; Pred. No. 0.0031;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCGV 15  
|||||  
Db 823 GTPGPGIAGGRCGV 837

## RESULT 9

US-10-301-822-28

; Sequence 28, Application US/10301822

; Publication No. US20030148410A1

; GENERAL INFORMATION:

; APPLICANT: Millennium Pharmaceuticals, Inc.

; APPLICANT: Berger, Allison

; APPLICANT: Guillemette, Tracy L.

; APPLICANT: Kamatkar, Shubhangi

; APPLICANT: Schlegel, Robert

; APPLICANT: Monahan, John E.

; APPLICANT: Thibodeau, Stephen N.

; APPLICANT: BURGART, Lawrence J.

; TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND

;; TITLE OF INVENTION: METHODS FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND  
;; FILE REFERENCE: MPM01-029P2RNM  
;; CURRENT APPLICATION NUMBER: US/10/301,822  
;; CURRENT FILING DATE: 2002-11-21  
;; PRIOR APPLICATION NUMBER: US 60/339,971  
;; PRIOR FILING DATE: 2001-12-10  
;; PRIOR APPLICATION NUMBER: US 60/361,978  
;; PRIOR FILING DATE: 2002-03-05  
;; PRIOR APPLICATION NUMBER: US 60/381,988  
;; PRIOR FILING DATE: 2002-05-20  
;; NUMBER OF SEQ ID NOS: 228  
;; SOFTWARE: FastSeq for Windows Version 4.0  
;; SEQ ID NO 28  
;; LENGTH: 1464  
;; TYPE: PRT  
;; ORGANISM: Homo Sapiens  
US-10-301-822-28

Query Match 100.0%; Score 80; DB 12; Length 1464;  
Best Local Similarity 100.0%; Pred. No. 0.0034;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGVY 15  
|||||

DB 947 GTPGPGIAGRGVY 961

RESULT 10  
US-10-060-036-159  
;; Sequence 159, Application US/10060036  
;; Publication No. US20030073144A1  
;; GENERAL INFORMATION:  
;; APPLICANT: Benson, Darin R.  
;; APPLICANT: Kalos, Michael D.  
;; APPLICANT: Lodes, Michael J.  
;; APPLICANT: Persing, David H.  
;; APPLICANT: Hepler, William T.  
;; APPLICANT: Jiang, Yuqiu  
;; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY  
;; TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER  
;; FILE REFERENCE: 210121.566  
;; CURRENT APPLICATION NUMBER: US/10/060,036  
;; CURRENT FILING DATE: 2002-01-30  
;; NUMBER OF SEQ ID NOS: 4560  
;; SOFTWARE: FastSeq for Windows Version 4.0  
;; SEQ ID NO 159  
;; LENGTH: 1464  
;; TYPE: PRT  
;; ORGANISM: Homo sapiens  
US-10-060-036-159

Query Match 100.0%; Score 80; DB 15; Length 1464;  
Best Local Similarity 100.0%; Pred. No. 0.0034;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGVY 15  
|||||

DB 947 GTPGPGIAGRGVY 961

RESULT 11  
US-10-171-311-36  
;; Sequence 36, Application US/10171311  
;; Publication No. US20030087270A1  
;; GENERAL INFORMATION:  
;; APPLICANT: Schlegel, Robert  
;; APPLICANT: Chen, Yan  
;; APPLICANT: Zhao, Xumei  
;; APPLICANT: Monahan, John  
;; APPLICANT: Kamatkar, Shubhangi  
;; APPLICANT: Glatz, Karen  
;; APPLICANT: Gannavairapu, Manjula

;; APPLICANT: Hoersch, Sebastian  
;; TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR  
;; TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY  
;; TITLE OF INVENTION: OF CERVICAL CANCER  
;; FILE REFERENCE: MPT-035  
;; CURRENT APPLICATION NUMBER: US/10/171,311  
;; CURRENT FILING DATE: 2002-06-12  
;; PRIOR APPLICATION NUMBER: US 60/298,159  
;; PRIOR FILING DATE: 2001-06-13  
;; PRIOR APPLICATION NUMBER: US 60/298,155  
;; PRIOR FILING DATE: 2001-06-13  
;; PRIOR APPLICATION NUMBER: US 60/335,936  
;; PRIOR FILING DATE: 2001-11-14  
;; NUMBER OF SEQ ID NOS: 238  
;; SOFTWARE: FastSeq for Windows Version 4.0  
;; SEQ ID NO 36  
;; LENGTH: 1464  
;; TYPE: PRT  
;; ORGANISM: Homo sapiens  
US-10-171-311-36

Query Match 100.0%; Score 80; DB 15; Length 1464;  
Best Local Similarity 100.0%; Pred. No. 0.0034;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGVY 15  
|||||

DB 947 GTPGPGIAGRGVY 961

RESULT 12  
US-10-216-705-21  
;; Sequence 21, Application US/10216705  
;; Publication No. US20030096973A1  
;; GENERAL INFORMATION:  
;; APPLICANT: Meristem Therapeutics, S.A.  
;; TITLE OF INVENTION: Recombinant Collagens and Derived Proteins Produced by Plants,  
;; TITLE OF INVENTION: Obtaining Such and Their Uses  
;; FILE REFERENCE: 1149-3 DIV  
;; CURRENT APPLICATION NUMBER: US/10/216,705  
;; CURRENT FILING DATE: 2002-08-09  
;; PRIOR APPLICATION NUMBER: US 09/331,347  
;; PRIOR FILING DATE: 1999-08-17  
;; NUMBER OF SEQ ID NOS: 22  
;; SOFTWARE: PatentIn version 3.1  
;; SEQ ID NO 21  
;; LENGTH: 1464  
;; TYPE: PRT  
;; ORGANISM: Homo sapiens  
US-10-216-705-21

Query Match 100.0%; Score 80; DB 15; Length 1464;  
Best Local Similarity 100.0%; Pred. No. 0.0034;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGVY 15  
|||||

DB 947 GTPGPGIAGRGVY 961

RESULT 13  
US-10-149-352-2  
;; Sequence 2, Application US/10149352  
;; Publication No. US20030105050A1  
;; GENERAL INFORMATION:  
;; APPLICANT: Berl, Rajinder  
;; TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES  
;; FILE REFERENCE: 06275-254US1  
;; CURRENT APPLICATION NUMBER: US/10/149,352  
;; CURRENT FILING DATE: 2002-06-10  
;; PRIOR APPLICATION NUMBER: PCT/GB00/04741  
;; PRIOR FILING DATE: 2000-12-12  
;; PRIOR APPLICATION NUMBER: GB 9929487.8

;; PRIOR FILING DATE: 1999-12-15  
;; NUMBER OF SEQ ID NOS: 14  
;; SOFTWARE: PatentIn Ver. 4.0  
;; SEQ ID NO 2  
;; LENGTH: 1464  
;; TYPE: PRT  
;; ORGANISM: Homo sapiens  
US-10-149-352-2

Query Match 100.0%; Score 80; DB 15; Length 1464;  
Best Local Similarity 100.0%; Pred. No. 0.0034;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCVV 15  
DB 947 GTPGPGIAGGRCVV 961

## RESULT 14

US-10-177-293-65  
;; Sequence 65, Application US/10177293  
;; Publication No. US20030124128A1  
;; GENERAL INFORMATION:

;; APPLICANT: Lillie, James  
;; APPLICANT: Glatc, Karen  
;; APPLICANT: Zhao, Xumei  
;; APPLICANT: Gannavarpu, Manjula  
;; APPLICANT: Kamatkar, Shubhangl  
;; APPLICANT: Mertens, Maureen  
;; APPLICANT: Myer, Vic  
;; APPLICANT: Wang, Youzhen  
;; APPLICANT: Xu, Yongyao  
;; APPLICANT: Hoersch, Sebastian  
;; APPLICANT: Monahan, John  
;; APPLICANT: Meyers, Rachel E.  
;; APPLICANT: Bast Jr., Robert C.  
;; APPLICANT: Hortobagyl, Gabriel N.  
;; APPLICANT: Pusztai, Lajos  
;; APPLICANT: Meric, Funda  
;; APPLICANT: Sahin, Aysegul  
;; APPLICANT: Mills, Gordon B.  
;; TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR IDENTIFICATION, ASSESSMENT,  
;; TITLE OF INVENTION: PREVENTION, AND THERAPY OF BREAST CANCER  
;; FILE REFERENCE: MRI-038  
;; CURRENT APPLICATION NUMBER: US/10/177,293  
;; CURRENT FILING DATE: 2002-06-21  
;; PRIOR APPLICATION NUMBER: US 60/299,887  
;; PRIOR FILING DATE: 2001-06-21  
;; PRIOR APPLICATION NUMBER: US 60/301,572  
;; PRIOR FILING DATE: 2001-06-27  
;; PRIOR APPLICATION NUMBER: US 60/306,501  
;; PRIOR FILING DATE: 2001-07-18  
;; PRIOR APPLICATION NUMBER: US 60/325,002  
;; PRIOR FILING DATE: 2001-09-25  
;; PRIOR APPLICATION NUMBER: US 60/362,585  
;; PRIOR FILING DATE: 2002-03-05  
;; PRIOR APPLICATION NUMBER: US 60/xxx,xxx  
;; PRIOR FILING DATE: 2002-05-14  
;; NUMBER OF SEQ ID NOS: 506  
;; SOFTWARE: FastSeq for Windows Version 4.0  
;; SEQ ID NO 65  
;; LENGTH: 1464  
;; TYPE: PRT  
;; ORGANISM: Homo sapiens  
US-10-177-293-65

Query Match 100.0%; Score 80; DB 15; Length 1464;  
Best Local Similarity 100.0%; Pred. No. 0.0034;  
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCVV 15  
DB 947 GTPGPGIAGGRCVV 961

## RESULT 15

US-09-895-674-1  
;; Sequence 1, Application US/09895674  
;; Publication No. US20030021821A1  
;; GENERAL INFORMATION:  
;; APPLICANT: Fertala, Andrzej  
;; APPLICANT: Ko, Frank  
;; TITLE OF INVENTION: Collagen and Collagen-like Peptide Containing Polymeric  
;; TITLE OF INVENTION: Matrices  
;; FILE REFERENCE: DRE-0032  
;; CURRENT APPLICATION NUMBER: US/09/895,674  
;; CURRENT FILING DATE: 2001-06-28  
;; PRIOR APPLICATION NUMBER: PCT/US01/  
;; PRIOR FILING DATE: 2001-06-25  
;; PRIOR APPLICATION NUMBER: 60/ 214,034  
;; PRIOR FILING DATE: 2000-06-23  
;; NUMBER OF SEQ ID NOS: 1  
;; SOFTWARE: PatentIn Ver. 2.1  
;; SEQ ID NO 1  
;; LENGTH: 234  
;; TYPE: PRT  
;; ORGANISM: Homo sapiens  
US-09-895-674-1

Query Match 88.8%; Score 71; DB 11; Length 234;  
Best Local Similarity 80.0%; Pred. No. 0.011;  
Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 GTPGPGIAGGRCVV 15  
DB 67 GTPGPGIAGGRCVV 81

Search completed: August 29, 2003, 18:30:43  
Job time : 26 secs

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GenCore version 5.1.6  
Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: August 29, 2003, 18:20:35 ; Search time 40 Seconds  
(without alignments)  
36.063 Million cell updates/sec

Title: US-09-935-417-1

Perfect score: 80

Sequence: 1 GTPGPGIAGRGVY 15

Scoring table:

BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 283308 seqs, 96168682 residues

Total number of hits satisfying chosen parameters: 283308

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%

Listing first 45 summaries

Database :

1: pir1:\*  
2: pir2:\*  
3: pir3:\*  
4: pir4:\*

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	80	100.0	779	1 CGB01S	collagen alpha 1(I)
2	80	100.0	1042	1 CGCH1S	collagen alpha 1(I)
3	80	100.0	1453	2 S21626	collagen alpha 1(I)
4	80	100.0	1464	1 CGHU1S	collagen alpha 1(I)
5	71	88.8	1418	2 T45467	collagen alpha 1(I)
6	71	88.8	1419	2 T41182	collagen alpha 1(I)
7	71	88.8	1487	1 CGHU6C	collagen alpha 1(I)
8	71	88.8	1487	2 B41182	collagen alpha 1(I)
9	68	85.0	1486	1 B40333	collagen alpha 1(I)
10	62	77.5	1763	2 S16366	collagen alpha 2(I)
11	59	73.8	618	2 S32436	collagen alpha 2(I)
12	58	72.5	290	2 A32249	collagen alpha 2(I)
13	58	72.5	1496	1 CGHU2V	collagen alpha 2(V)
14	58	72.5	1497	2 I49607	procollagen type V
15	57	71.2	310	2 I50696	collagen alpha 1(I)
16	57	71.2	636	2 S41067	collagen alpha 1(I)
17	57	71.2	886	2 I50694	collagen alpha 1(I)
18	57	71.2	1049	1 CGB07S	collagen alpha 1(I)
19	57	71.2	1414	1 S23809	collagen alpha 1(I)
20	57	71.2	1464	2 S59856	collagen alpha 1(I)
21	57	71.2	1466	1 CGHU7L	collagen alpha 1(I)
22	56	70.0	677	2 S23296	collagen alpha 2(I)
23	56	70.0	920	2 A45748	collagen alpha 1(V)
24	56	70.0	1366	1 CGHU2S	collagen alpha 2(V)
25	56	70.0	1669	1 CGHU4B	collagen alpha 1(I)
26	55	68.8	1669	1 CGMS4B	collagen alpha 1(I)
27	55	68.8	1688	2 A53330	collagen alpha 2(I)
28	55	68.8	1051	2 A35763	collagen alpha 2(C)
29	55	68.8	1492	2 A40333	collagen alpha 1(I)

30	54	67.5	298	2 T32371	hypothetical prote
31	53	66.2	266	2 T22706	hypothetical prote
32	53	66.2	287	2 T15779	hypothetical prote
33	53	66.2	299	2 T22705	hypothetical prote
34	53	66.2	305	2 T30165	hypothetical prote
35	53	66.2	307	2 T19582	hypothetical prote
36	53	66.2	673	1 CGHU6C	collagen alpha 1(I)
37	53	66.2	680	1 CGHU1D	collagen alpha 1(X)
38	53	66.2	751	2 S64741	cuticle collagen
39	53	66.2	775	2 A61228	collagen alpha 2(I)
40	53	66.2	959	2 S32605	collagen alpha 3(V)
41	53	66.2	1142	2 JX0369	collagen alpha 1(X)
42	53	66.2	1603	2 S23810	collagen alpha 1(X)
43	53	66.2	1752	2 A45407	collagen alpha 3(I)
44	53	66.2	1758	2 T29350	hypothetical prote
45	53	66.2	1759	2 T29351	collagen alpha 2(I)

#### ALIGNMENTS

##### RESULT 1

CGB01S

collagen alpha 1(I) chain - bovine (tentative sequence) (fragments)

C:Species: Bos primigenius taurus (cattle)

C:Date: 24-Apr-1984 #sequence\_revision 31-Dec-1993 #text\_change 31-Mar-2000

C:Accession: A91193; A91229; A91387; A91211; A91201; A91200; A43048; A02853

R:Rauterberg, J.; Timpl, R.; Furthmayr, H.

Eur. J. Biochem. 27, 231-237, 1972

A:Title: Structural characterization of N-terminal antigenic determinants in calf and

A:Reference number: A91193; MUID:7225534; PMID:4115172

A:Accession: A91193

A:Molecule type: protein

A:Residues: 1-19 <RAU>

A:Experimental source: skin

A>Note: the epsilon carbon of Lys-9, by homology with the rat alpha 1(I) chain, is co

R:Rietzek, P.P.; Kuehn, K.

Eur. J. Biochem. 52, 77-82, 1975

A:Title: The covalent structure of collagen: amino-acid sequence of the cyanogen-brom

A:Reference number: A91229; MUID:76022320; PMID:1164916

A:Accession: A91229

A:Molecule type: protein

A:Residues: 20-145 <RTD>

A:Experimental source: skin

A>Note: Lys-103 is hydroxylated and binds glucosylgalactose

R:Rietzek, P.P.; Wendt, P.; Kell, I.; Kuehn, K.

FEBS Lett. 26, 74-76, 1972

A:Title: The covalent structure of collagen: amino acid sequence of alpha1-CB3 from c

A:Reference number: A91387; MUID:73049499; PMID:4673951

A:Accession: A91387

A:Molecule type: protein

A:Residues: 146-294 <RT2>

A:Experimental source: skin

R:Rietzek, P.P.; Rexrodt, F.W.; Hopper, K.E.; Kuehn, K.

Eur. J. Biochem. 38, 396-400, 1973

A:Title: The covalent structure of collagen. 2. The amino-acid sequence of alpha1-CB7

A:Reference number: A91211; MUID:74086118; PMID:4359390

A:Accession: A91211

A:Molecule type: protein

A:Residues: 295-562 <RT3>

A:Experimental source: skin

R:Wendt, P.; Mark, K.V.D.; Rexrodt, F.; Kuehn, K.

Eur. J. Biochem. 30, 169-183, 1972

A:Title: The covalent structure of collagen. The amino-acid sequence of the 112 resid

A:Reference number: A91201; MUID:73042276; PMID:4343808

A:Accession: A91201

A:Molecule type: protein

A:Residues: 563-675 <WEN>

A:Experimental source: skin

R:Rietzek, P.P.; Rexrodt, F.W.; Wendt, P.; Stark, M.; Kuehn, K.

Eur. J. Biochem. 30, 163-168, 1972

A:Title: The covalent structure of collagen. Amino acid sequence of peptide alpha1-CB

A:Reference number: A91200; MUID:73042275; PMID:4343807

A:Accession: A91200  
 A:Molecule type: protein  
 A:Residues: 676-758 <FI4>  
 A:Experimental source: skin  
 A:Note: Pro-726 is the only 3-hydroxyproline and the only hydroxylated proline in position 726.  
 R:Rautenberg, J.; Fietzek, F.; Rexrodt, F.; Becker, U.; Stark, M.; Kuehn, K.  
 FEBS Lett. 21, 75-79, 1972  
 A:Title: The amino acid sequence of the carboxyterminal nonhelical cross link region of A:Reference number: A43048  
 A:Accession: A43048  
 A:Molecule type: protein  
 A:Residues: 759-779 <RA2>  
 A:Experimental source: skin  
 C:Comment: Lysines at positions 115, 124, 274, 346, 424, 496, 658, and 670 may be hydroxylated.  
 C:Comment: Prolines in the third position of the tripeptide repeating unit (G-X-Y) are hydroxylated.  
 C:Comment: The order of the eight CNBr peptides in the alpha 1(I) chain of bovine skin is 9, 149, 268, and 217 residues.  
 C:Comment: The complete chain contains 1052 residues.  
 C:Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homology;  
 C:Keywords: collid coll; extracellular matrix; glycoprotein; pyroglutamic acid; trimer;  
 F:1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental

Query Match 100.0%; Score 80; DB 1; Length 779;  
 Best Local Similarity 100.0%; Pred. No. 7.5e-05;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 GTPGPGIAGRGVY 15  
 |||||  
 Db 509 GTPGPGIAGRGVY 523

RESULT 2  
 CGCH15  
 Collagen alpha 1(I) chain - chicken (tentative sequence) (fragments)  
 C:Species: Gallus gallus (chicken)  
 C:Date: 12-Aug-1991 #sequence revision 06-Jul-1982 #text-change 31-Mar-2000  
 C:Accession: A90458; A90181; A02857  
 R:Highberger, J.H.; Corbett, C.; Dixit, S.N.; Yu, W.; Seyer, J.M.; Kang, A.H.; Gross, J.  
 Biochemistry 21, 2048-2055, 1982  
 A:Title: Amino acid sequence of chick skin collagen alpha 1(I)-C88 and the complete prime  
 A:Reference number: A90458; MUID:82231995; PMID:7093229  
 A:Accession: A90458  
 A:Molecule type: protein  
 A:Residues: 1-1036 <HIG>  
 A:Experimental source: skin  
 A:Note: This is the latest in a series of papers from these workers elucidating the sequence of the alpha 1(I) chain.  
 R:Exre, D.R.; Glimcher, M.J.  
 Biochem. Biophys. Res. Commun. 48, 720-726, 1972  
 A:Title: Evidence for a previously undetected sequence at the carboxyterminus of the alpha 1(I) chain.  
 A:Reference number: A90181; MUID:72243016; PMID:5047697  
 A:Accession: A90181  
 A:Molecule type: protein  
 A:Residues: 1037-1042 <EYR>  
 A:Experimental source: skin  
 C:Comment: Lysines at positions 103, 700, 934, and 946 above may be hydroxylated in some C:Comment: Most of the prolines at the third position of the tripeptide repeating unit (G-X-Y) are hydroxylated.  
 C:Comment: Pro-1002 is the only 3-hydroxyproline and the only hydroxylated proline in position 1002.  
 C:Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homology;  
 C:Keywords: collid coll; extracellular matrix; glycoprotein; pyroglutamic acid; trimer;  
 F:1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental

Query Match 100.0%; Score 80; DB 1; Length 1042;  
 Best Local Similarity 100.0%; Pred. No. 9.9e-05;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 GTPGPGIAGRGVY 15  
 |||||  
 Db 785 GTPGPGIAGRGVY 799

RESULT 3  
 521626

collagen alpha 1(I) chain precursor - mouse  
 C:Species: Mus musculus (house mouse)  
 C:Date: 13-Jan-1995 #sequence revision 25-Apr-1997 #text-change 13-Aug-1999  
 C:Accession: S57243; S16374; A23982; I49559; I45557; S39789; I48300; S21626  
 R:Li, S.W.; Khillan, J.; Prockop, D.J.  
 Matrix Biol. 14, 593-595, 1994  
 A:Title: The complete cDNA coding sequence for the mouse pro-alpha-1(I) chain of type I:Reference number: S57243  
 A:Accession: S57243  
 A:Molecule type: mRNA  
 A:Residues: 1-1453 <LIS>  
 A:Cross-references: EMBL:008020; NID:9470673; PIDN:AAA8912.1; PID:9470674  
 R:Mesaeranta, M.; Toman, D.; de Crombrughe, B.; Vuorio, E.  
 Biochim. Biophys. Acta 1089, 241-243, 1991  
 A:Title: Specific hybridization probes for mouse type I, II, III and IX collagen mRNA  
 A:Reference number: S16176; MUID:91274355; PMID:2054384  
 A:Accession: S16374  
 A:Status: preliminary  
 A:Molecule type: DNA  
 A:Residues: 1442-1453 <MET>  
 A:Cross-references: EMBL:X57981; NID:950484; PIDN:CAA1046.1; PID:950485  
 R:French, B.T.; Lee, W.H.; Maul, G.G.  
 Gene 39, 311-312, 1985  
 A:Title: Nucleotide sequence of a cDNA clone for mouse proalpha1(I) collagen protein.  
 A:Reference number: A23982; MUID:86137403; PMID:3841523  
 A:Accession: A23982  
 A:Molecule type: mRNA  
 A:Residues: 518-1128 <FRE>  
 A:Cross-references: GB:M14423; NID:9192261; PIDN:AAA3733.1; PID:9192262  
 R:Monson, J.M.; Friedman, J.; McCarthy, B.J.  
 Mol. Cell. Biol. 2, 1362-1371, 1982  
 A:Title: DNA sequence analysis of a mouse pro-alpha-1(I) procollagen gene: Evidence for A:Reference number: I49559; MUID:83141374; PMID:6288597  
 A:Accession: I49559  
 A:Status: preliminary; translated from GB/EMBL/DBJ  
 A:Molecule type: DNA  
 A:Residues: 735-1130 <RES>  
 A:Cross-references: GB:M17491; NID:9192263; PIDN:AAA37334.1; PID:9192264  
 R:Harbers, K.; Kuehn, M.; Delius, H.; Jaenisch, R.  
 Proc. Natl. Acad. Sci. U.S.A. 81, 1504-1508, 1984  
 A:Title: Insertion of retrovirus into the first intron of alpha1(I) collagen gene leads A:Reference number: I49557; MUID:86170331; PMID:6324198  
 A:Accession: I49557  
 A:Status: preliminary; translated from GB/EMBL/DBJ  
 A:Molecule type: DNA  
 A:Residues: 1-25 <RNE>  
 A:Cross-references: GB:K01688; NID:9192246; PIDN:AAA37330.1; PID:9553881  
 R:Penton, S.P.; Lamande, S.R.; Hannagan, M.; Stacey, A.; Jaenisch, R.; Bateman, J.F.  
 Biochim. Biophys. Acta 1216, 469-474, 1993  
 A:Title: Genomic sequence of mouse COL1A1 encoding the collagen propeptides.  
 A:Reference number: S39789; MUID:94092741; PMID:8268229  
 A:Accession: S39789  
 A:Molecule type: DNA  
 A:Residues: 1-80, 'E', 82-105, 'D', 107-147 <REF>  
 A:Cross-references: EMBL:X54876; NID:950486; PIDN:CAA38657.1; PID:950487  
 A:Accession: X54876  
 A:Molecule type: DNA  
 A:Residues: 1-80, 'E', 82-105, 'D', 107-147 <REF>  
 A:Cross-references: EMBL:X54876; NID:950486; PIDN:CAA38657.1; PID:950487  
 A:Accession: X54876  
 A:Gene: COL1A1  
 A:Introns: 770/3; 788/3; 806/3; 842/3; 860/3; 878/3; 932/3; 968/3; 1004/3; 1022/3; 10 C:Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homology  
 C:Keywords: collid coll; extracellular matrix; glycoprotein; heterotrimer; triple helix  
 F:1-22/Domain: signal sequence #status predicted <SIG>  
 F:23-151/Domain: amino-terminal propeptide #status predicted <PRO>  
 F:30-89/Domain: von Willebrand factor type C repeat homology <WVC>  
 F:152-1453/Product: collagen alpha 1(I) chain #status predicted <MAT>  
 F:1224-1453/Domain: fibrillar collagen carboxyl-terminal homology <FCG>

Query Match 100.0%; Score 80; DB 2; Length 1453;  
 Best Local Similarity 100.0%; Pred. No. 0 00014;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GTPGPGIAGORGVV 15  
 |||||  
 Db 936 GTPGPGIAGORGVV 950

RESULT 4  
 CGHUIS  
 collagen alpha 1(I) chain precursor - human  
 N:Alternate names: procollagen alpha 1(I) chain  
 C:Species: Homo sapiens (man)  
 C:Date: 12-Aug-1981 #sequence\_revision 04-Oct-1996 #text\_change 31-Dec-2000  
 C:Accession: 160114; S01143; A93335; I55254; A39943; I55237; A35233; S09400; B90567; S115269; A29439; I53466; A02852; I37247  
 R:D'Alessio, M.; Bernard, M.; Pretorius, P.J.; de Wet, W.; Ramirez, F.; Pretorius, P.J.  
 Gene 67, 105-115, 1988

A:Title: Complete nucleotide sequence of the region encompassing the first twenty-five  
 A:Reference number: 160114; MUID:88329734; PMID:2843432  
 A:Accession: 160114

A:Status: translated from GB/EMBL/DBJ  
 A:Molecule type: DNA  
 A:Residues: 1-369, 'L', 371-589 <DAL>  
 A:Cross-references: GB:M20789; NID:9179593; PIDN:AA59373.1; PID:9179594  
 R:Tromp, G.; Kuvshinov, H.; Stacey, A.; Shikata, H.; Baldwin, C.T.; Jaenisch, R.; Prockop  
 Blochem. J. 253, 919-922, 1988

A:Title: Structure of a full-length cDNA clone for the prepro-alpha-1(I) chain of human  
 A:Reference number: S01143; MUID:89025644; PMID:318743  
 A:Accession: S01143

A:Molecule type: mRNA  
 A:Residues: 1-472 <TR>  
 A:Cross-references: EMBL:X07884; NID:930015; PIDN:CAA30731.1; PID:930016; GB:M36546; NID  
 A:Note: submitted to the EMBL/GenBank/DBJ databases by Prockop, D.J., 13-JUN-1988

R:Chu, M.L.; de Wet, W.; Bernard, M.; Ding, J.F.; Morabito, M.; Myers, J.; Williams, C.;  
 Nature 310, 337-340, 1984

A:Title: Human proalpha1(I) collagen gene structure reveals evolutionary conservation of  
 A:Reference number: A93335; MUID:84270697; PMID:6462220  
 A:Accession: A93335

A:Molecule type: DNA  
 A:Residues: 1-58, 'Q', 60-181 <CHD>  
 A:Cross-references: EMBL:X00820; NID:935657; PIDN:CAA52594.1; PID:935658  
 R:Rosouw, C.M.S.; Vergeer, W.P.; du Plooy, S.J.; Bernard, M.P.; Ramirez, F.; de Wet, W.  
 J. Biol. Chem. 262, 15151-15157, 1987

A:Title: DNA sequences in the first intron of the human pro-alpha 1(I) collagen gene en  
 A:Reference number: I55254; MUID:88033098; PMID:2822714  
 A:Accession: I55254

A:Status: translation not shown; translated from GB/EMBL/DBJ  
 A:Molecule type: DNA  
 A:Residues: 1-45 <ROS>  
 A:Cross-references: GB:J02829; NID:9180387; PIDN:AA51993.1; PID:9180388  
 R:Bornstein, P.; McKay, J.; Morishima, J.K.; Devareyalu, S.; Gellinas, R.E.  
 Proc. Natl. Acad. Sci. U.S.A. 84, 8869-8873, 1987

A:Title: Regulatory elements in the first intron contribute to transcriptional control c  
 A:Reference number: A39943; MUID:88097389; PMID:3480516  
 A:Accession: A39943

A:Molecule type: DNA  
 A:Residues: 1-34 <BOR>  
 A:Cross-references: GB:J03559; NID:9180876; PIDN:AA52052.1; PID:955238  
 R:Chu, M.L.; de Wet, W.; Bernard, M.; Ramirez, F.  
 J. Biol. Chem. 260, 2315-2320, 1985

A:Title: Fine structural analysis of the human pro-alpha 1 (I) collagen gene. Promoter s  
 A:Reference number: I55237; MUID:85130970; PMID:2857713  
 A:Accession: I55237

A:Status: translation not shown; translated from GB/EMBL/DBJ  
 A:Molecule type: DNA  
 A:Residues: 1-34 <CH2>  
 A:Cross-references: GB:M10627; NID:9180383; PIDN:AA51992.1; PID:9553226  
 R:Wirtz, M.R.; Keene, D.R.; Horl, R.; Glanville, R.W.; Steinmann, B.; Rao, V.H.; Hollist  
 J. Biol. Chem. 265, 6312-6317, 1990

A:Title: In vivo and in vitro noncovalent association of excised alpha1(I) amino-termin

rome, type VII.  
 A:Reference number: A35233; MUID:90202908; PMID:2318855  
 A:Accession: A35233

A:Molecule type: protein  
 A:Residues: 33-52 <MR>  
 A:Note: this propeptide fragment remained non-covalently bound to a defective, unclea  
 R:Wells, D.; d'Alessio, M.; Ramirez, F.; de Wet, W.; Cole, W.G.; Chan, D.; Bateman, J.  
 EMBO J. 8, 1705-1710, 1989

A:Title: A base substitution in the exon of a collagen gene causes alternative splici  
 A:Reference number: S09400; MUID:89356643; PMID:2767050  
 A:Accession: S09400

A:Molecule type: mRNA  
 A:Residues: 156-183 <MEI>  
 R:Click, E.M.; Bornstein, P.  
 Biochemistry 9, 4699-4706, 1970

A:Title: Isolation and characterization of the cyanogen bromide peptides from the alp  
 A:Reference number: A90567; MUID:71038625; PMID:5529814  
 A:Contents: CNBR0-1, CNBR2, CNBR4, CNBR5

A:Accession: B90567

A:Molecule type: protein  
 A:Residues: 162-198, 'Z', 200-201, 'Z', 203-206, 'Z', 208-209, 'Z', 211-228, 'B', 230, 'BB', 233,  
 A:Experimental source: skin  
 A:Note: evidence for 170-allysine

R:Baetge, B.; Notbohm, H.; Diebold, J.; Lehmann, H.; Bodo, M.; Deutzmann, R.; Mueller  
 Eur. J. Biochem. 192, 153-159, 1990

A:Title: A critical crosslink region in human-bone-derived collagen type I. Specific  
 A:Reference number: S11372; MUID:90382436; PMID:2169412  
 A:Accession: S11372

A:Molecule type: protein  
 A:Residues: 175-187, 274-287, 'P', 289 <BAE>  
 A:Note: sequence of collagen alpha 1(S)(I) isolated from bone after pepsin digestion

R:Deak, S.B.; Scholz, P.M.; Amenta, P.S.; Constantinou, C.D.; Levi-Minzl, S.A.; Gonzal  
 J. Biol. Chem. 266, 21827-21832, 1991

A:Title: The substitution of arginine for glycine 85 of the alpha 1(I) procollagen ch  
 A:Reference number: I55342; MUID:92042092; PMID:1718984  
 A:Accession: I55342

A:Status: translated from GB/EMBL/DBJ  
 A:Molecule type: mRNA  
 A:Residues: 258-268, 11347-1357 <DEA>  
 A:Cross-references: GB:S67495; NID:9239007; PIDN:AA820350.1; PID:9239008

A:Note: sequences from the 5' and 3' ends only are shown; mutant sequence 263-Arg rep  
 R:Morgan, P.H.; Jacobs, H.G.; Segrest, J.P.; Cunningham, L.W.  
 J. Biol. Chem. 245, 5042-5048, 1970

A:Title: Comparative study of glycopeptides derived from selected vertebrate collagen  
 A:Reference number: A92069; MUID:71001508; PMID:4319110  
 A:Accession: A92069

A:Molecule type: protein  
 A:Residues: 263-268 <MOR>  
 A:Experimental source: skin  
 A:Note: attachment of 2-O-alpha-D-glucosyl-O-beta-D-galactose to 5-hydroxylysine

R:Labhard, M.E.; Hollister, D.W.  
 Matrix 10, 124-130, 1990

A:Title: Segmental amplification of the entire helical and telopeptide regions of the  
 A:Reference number: S15989; MUID:90326017; PMID:2374517  
 A:Accession: S15989

A:Molecule type: mRNA  
 A:Residues: 281-302, 402-420, 823-843, 925-944, 1026-1045, 1143-1162 <LAB>  
 R:Wirtz, M.R.; Rao, V.H.; Glanville, R.W.; Labhard, M.E.; Pretorius, P.J.; de Vries,  
 Connect. Tissue Res. 29, 1-11, 1993

A:Title: A cysteine for glycine substitution at position 175 in an alpha 1 (I) chain  
 A:Reference number: I52905; MUID:93339042; PMID:8339541  
 A:Accession: I52905

A:Status: translated from GB/EMBL/DBJ  
 A:Molecule type: mRNA  
 A:Residues: 342-352, 'C', 354-359 <MI2>  
 A:Cross-references: GB:S64717; NID:9408195; PIDN:AA827677.1; PID:9408196

A:Note: mutant sequence from patient with osteogenesis imperfecta  
 R:Bernard, M.P.; Chu, M.L.; Myers, J.C.; Ramirez, F.; Elkenberry, E.F.; Prockop, D.J.  
 Biochemistry 22, 5213-5223, 1983

A:Title: Nucleotide sequences of complementary deoxyribonucleic acids for the proalph  
 A:Reference number: A90476; MUID:84080385; PMID:6689127  
 A:Accession: A90476

A:Molecule type: mRNA  
 A:Residues: 425-1250, 'X', 1252-1328, 'S', 1330-1390, 'X', 1392-1464 <BER>  
 A:Cross-references: GB:K01228; NID:G180391; PIDN:AAA51995.1; PID:G180392  
 A:Note: sequence partially completed for missing nucleotides by A29439  
 R:Chu, M.L.; Garguilo, V.; Williams, C.J.; Ramirez, F.  
 J. Biol. Chem. 260, 691-694, 1985  
 A:Title: Multixon deletion in an osteogenesis imperfecta variant with increased type II  
 A:Reference number: A22161; MUID:85104934; PMID:2981843  
 A:Accession: A22161  
 A:Molecule type: DNA  
 A:Residues: 472-594, 'R', 596-607 <CH3>  
 A:Cross-references: GB:K03178; GB:K03179; NID:G179612; NID:G179613; PIDN:AAA51847.1; PID  
 A:Note: the authors translated the codon CGT for residue 595 as Pro  
 R:Wallis, G.A.; Starman, B.J.; Zinn, A.B.; Byers, P.H.  
 Am. J. Hum. Genet. 46, 1034-1040, 1990  
 A:Title: Variable expression of osteogenesis imperfecta in a nuclear family is explained  
 A:Reference number: A35336; MUID:90252792; PMID:2339700  
 A:Accession: A35336  
 A:Molecule type: mRNA  
 A:Residues: 710-720, 'E', 722-737, 'E', 739-745 <MAL>  
 A:Note: the authors translated the codons CAG for 721 and CGT for 738 as Glu  
 R:Forlino, A.; Zolietzi, F.; Valli, M.; Pignatelli, P.F.; Cetta, G.; Brunelli, P.C.; Motte  
 Hum. Mol. Genet. 3, 2201-2206, 1994  
 A:Title: Severe (type II) osteogenesis imperfecta due to glycine substitutions in the c  
 A:Reference number: I54365; MUID:95187161; PMID:7881420  
 A:Accession: I54365  
 A:Molecule type: DNA  
 A:Status: translated from GB/EMBL/DBJ  
 A:Residues: 746-766, 'S', 768-781 <FOR>  
 A:Cross-references: GB:I47667; NID:G1009093; PIDN:AA859576.1; PID:G1009094  
 R:Chesser, S.D.; Wallis, G.A.; Byers, P.H.  
 J. Biol. Chem. 268, 18218-18225, 1993  
 A:Title: Mutations in the carboxyl-terminal propeptide of the pro alpha 1(I) chain of ty  
 A:Reference number: A47426; MUID:93352646; PMID:8349697  
 A:Accession: A47426  
 A:Molecule type: mRNA  
 A:Residues: 1179-1276, 'H', 1278-1336, 1339-1387, 'R', 1389-1464 <CHE>  
 A:Cross-references: GB:564556; NID:G407589; PIDN:AA827856.1; PID:G407590  
 A:Note: sequence extracted from NCBI backbone (NCBIN:136444, NCBI:136445)  
 A:Note: does not represent an experimentally determined sequence but three different mut  
 A:Accession: B47426  
 A:Molecule type: mRNA  
 A:Residues: 1179-1464 <CH4>  
 A:Experimental source: normal dermal fibroblast culture  
 A:Accession: C47426  
 A:Molecule type: mRNA  
 A:Residues: 1179-1276, 'H', 1278-1464 <CH5>  
 A:Experimental source: fetal cell 86-237  
 A:Accession: D47426  
 A:Molecule type: mRNA  
 A:Residues: 1179-1336, 1339-1464 <CH6>  
 A:Experimental source: fetal cell 86-146  
 A:Accession: E47426  
 A:Molecule type: mRNA  
 A:Residues: 1179-1387, 'R', 1389-1464 <CH7>  
 A:Experimental source: fetal cell 88-251  
 R:Chen, D.H.; Apore, S.; Eyre, D.R.; Starman, B.J.; Andreasen, P.; Charbonneau, H.; Nid  
 J. Biol. Chem. 263, 14605-14607, 1988  
 A:Title: Substitution of Cysteine for Glycine within the Carboxyl-terminal Telopeptide c  
 A:Reference number: I55269; MUID:89008319; PMID:3170557  
 A:Accession: I55269  
 A:Status: translated from GB/EMBL/DBJ  
 A:Molecule type: DNA  
 A:Residues: 1187-1194, 'C', 1196-1220 <CON>  
 A:Cross-references: GB:M23213; NID:G340842; PIDN:AA859363.1; PID:G499622  
 A:Note: mutant sequence from a patient with mild osteogenesis imperfecta  
 R:Meekelae, J.K.; Raassina, M.; Virta, A.; Vuorio, E.  
 Nucleic Acids Res. 16, 349, 1988  
 A:Title: Human pro-alpha-1(I) collagen: cDNA sequence for the C-propeptide domain.

Query Match 100.0%; Score 80; DB 1; Length 1464;  
 Best Local Similarity 100.0%; Pred. No. 0.00014;

Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 Oy 1 GTPPGGIAGORGV 15  
 Db 947 GTPPGGIAGORGV 961

## RESULT 5

T45467

collagen alpha 1(II) chain precursor [imported] - horse

N:Alternate names: type II collagen

C:Species: Equus caballus (domestic horse)

C:Date: 31-Jan-2000 #sequence\_revision 31-Jan-2000 #text\_change 04-Mar-2000

C:Accession: T45467

R:Richardson, D.W.; Dodge, G.R.

submitted to the EMBL Data Library, June 1996

A:Description: Cloning of equine type II collagen and modulation of its expression in

A:Reference number: 222977

A:Accession: T45467

A:Status: preliminary; translated from GB/EMBL/DBJ

A:Molecule type: mRNA

A:Residues: 1-1418 &lt;RIC&gt;

A:Cross-references: EMBL:U62528; PIDN:AA805773.1

C:Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homolo

## Query Match

88.8%; Score 71; DB 2; Length 1418;

Best Local Similarity 80.0%; Pred. No. 0.0036;

Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Oy 1 GTPPGGIAGORGV 15

Db 900 GTPPGGIAGORGV 914

## RESULT 6

A41182

collagen alpha 1(II) chain precursor - mouse

C:Species: Mus musculus (house mouse)

C:Date: 28-May-1992 #sequence\_revision 28-May-1992 #text\_change 13-Aug-1999

C:Accession: A41182; A44885

R:Meisner, M.; Roman, D.; de Crombrughe, B.; Vuorio, E.

J. Biol. Chem. 266, 16862-16867, 1991

A:Title: Mouse type II collagen gene. Complete nucleotide sequence, exon structure, a

A:Reference number: A41182; MUID:91358489; PMID:1885613

A:Accession: A41182

A:Status: preliminary; not compared with conceptual translation

A:Molecule type: DNA

A:Residues: 1-1419 &lt;MET&gt;

A:Cross-references: GB:M65161

R:Cheah, K.S.; Lau, E.T.; Au, P.K.; Tam, P.P.

Development 111, 945-953, 1991

A:Title: Expression of the mouse alpha 1(II) collagen gene is not restricted to carti

A:Reference number: A44885; MUID:91347939; PMID:1879363

A:Accession: A44885

A:Molecule type: DNA

A:Residues: 1-28 &lt;CHE&gt;

A:Cross-references: GB:563190; NID:G234368; PIDN:AA819627.1; PID:G234369

A:Note: sequence extracted from NCBI backbone (NCBIN:63190, NCBI:63192)

C:Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homolo

C:Keywords: alternative splicing; coiled coil; extracellular matrix; glycoprotein; tr

F;1191-1419/Domains: fibrillar collagen carboxyl-terminal homology &lt;FC&gt;

## Query Match

88.8%; Score 71; DB 2; Length 1419;

Best Local Similarity 80.0%; Pred. No. 0.0036;

Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Oy 1 GTPPGGIAGORGV 15

Db 901 GTPPGGIAGORGV 915

## RESULT 7

CGH06C

collagen alpha 1(II) chain precursor [validated] - human  
 N.Alternate names: chondrocalcin, collagen alpha 1(II) chain  
 C.Species: Homo sapiens (man)  
 C.Date: 28-May-1986 #sequence, revision 01-Sep-1995 #text, change 08-Dec-2000  
 C.Accession: A38513; S66715; S24270; A24828; S06496; A35428; A30147; A33116; S64674; S63750; I37251; I37252; I37253; I37254; I55338; I59535; I61910  
 R.Ryan, M.C.; Sieraski, M.; Sandell, L.J.  
 Genomics 8, 41-48, 1990  
 A.Title: The human type II procollagen gene: identification of an additional protein-cod  
 A.Reference number: A38513; MUID:91184811; PMID:2081599  
 A.Accession: A38513  
 A.Molecule type: DNA  
 A.Residues: 1-103 <RYA>  
 A.Cross-references: GB:M20299; NID:9180883; PIDN:AAA73873.1; PID:9180884  
 R.Su, M.W.; Lee, B.; Ramirez, F.; Machado, M.; Horton, W.  
 Nucleic Acids Res. 17, 9473, 1989  
 A.Title: Nucleotide sequence of the full length cDNA encoding for human type II procolla  
 A.Reference number: S06715; MUID:90067946; PMID:2587267  
 A.Accession: S06715  
 A.Molecule type: mRNA  
 A.Residues: 1-28, 'R', '99-1487 <SU2>  
 A.Cross-references: EMBL:X16468; NID:929515; PIDN:CAA34488.1; PID:929516  
 A.Note: alternative splice form 1  
 R.Vikkula, M.; Metsaeranta, M.; Syvaenen, A.C.; Ala-Kokko, L.; Vuorio, E.; Peltonen, L.  
 Blochem. J. 285, 287-294, 1992  
 A.Title: Structural analysis of the regulatory elements of the type-II procollagen gene.  
 A.Reference number: S24270; MUID:92344585; PMID:1637314  
 A.Status: translation not shown  
 A.Molecule type: DNA  
 A.Residues: 1-28 <VIK>  
 A.Cross-references: EMBL:X58709; GB:S40537; NID:935659  
 A.Note: This translation is not annotated in Genbank entry HSPROCOE1, release 111.0  
 R.Nunez, A.M.; Kohno, K.; Martin, G.R.; Yamada, Y.  
 Gene 44, 11-16, 1986  
 A.Title: Promoter region of the human pro-alpha-1(II)-collagen gene.  
 A.Reference number: A24828; MUID:87031574; PMID:3021582  
 A.Accession: A24828  
 A.Molecule type: DNA  
 A.Residues: 1-8, 'T', '10-28 <NUN>  
 A.Cross-references: GB:M25698; NID:9180872; PIDN:AAA52051.1; PID:9553237  
 R.Baldwin, C.T.; Reginaldo, A.M.; Smith, C.; Jimenez, S.A.; Prockop, D.J.  
 Blochem. J. 262, 521-528, 1989  
 A.Title: Structure of cDNA clones coding for human type II procollagen. The alpha-1(II)  
 A.Reference number: S06496; MUID:90026318; PMID:2803268  
 A.Accession: S06496  
 A.Molecule type: mRNA  
 A.Residues: 7-28, 'R', '99-157, 'P', '159-440, 'G', '442-456, 'E', '458-640, 'A', '642-831, 'PA', '834, 'F'  
 A.Cross-references: EMBL:X16711; NID:930040; PIDN:CAA34683.1; PID:930041  
 A.Note: alternative splice form 1  
 R.Ryan, M.C.; Sandell, L.J.  
 J. Biol. Chem. 265, 10334-10339, 1990  
 A.Title: Differential expression of a cysteine-rich domain in the amino-terminal propept  
 A.Reference number: A35428; MUID:90285153; PMID:2355003  
 A.Accession: A35428  
 A.Status: not compared with conceptual translation  
 A.Molecule type: mRNA  
 A.Residues: 27-81, 'L', '83-103 <RYA2>  
 A.Note: alternative splice form 2; splicing appears to be under developmental regulation  
 R.Su, M.W.; Benson-Chanda, V.; Vissing, H.; Ramirez, F.  
 Genomics 4, 438-441, 1989  
 A.Title: Organization of the exons coding for Pro alpha-1(II) collagen N-propeptide cont  
 A.Reference number: A30147; MUID:89233138; PMID:2714801  
 A.Accession: A30147  
 A.Molecule type: DNA  
 A.Residues: 104-157, 'P', '159-236 <SUN>  
 A.Cross-references: GB:J03065; GB:M23660; GB:M25655; GB:M25656; GB:M25730; GB:M2168; GE  
 R.Ala-Kokko, L.; Baldwin, C.T.; Moskowitz, R.W.; Prockop, D.J.  
 Proc. Natl. Acad. Sci. U.S.A. 87, 6565-6568, 1990  
 A>Title: Single base mutation in the type II procollagen gene (COL2A1) as a cause of pr  
 A.Reference number: A94227; MUID:90370826; PMID:1975693  
 A.Accession: A33116

A.Molecule type: DNA  
 A.Residues: 171-172, 'C', '174-175 <ALA>  
 A>Note: mutant sequence from a family with primary generalized osteoartru  
 R.Diab, M.; Wu, J.J.; Eyre, D.R.  
 Biochem. J. 314, 327-332, 1996  
 A.Title: Collagen type IX from human cartilage: a structural profile of Intermolecula  
 A.Reference number: S64673; MUID:96195147; PMID:8660302  
 A.Accession: S64674  
 A.Molecule type: protein  
 A.Residues: 188-189, 'X', '191-195; 1224-1230, 'X', '1232-1236 <DIA>  
 R.Franc, S.; Marzin, E.; Boutillon, M.M.; Lafont, R.; Lechene de la Porte, P.; Herbag  
 Eur. J. Biochem. 234, 125-131, 1995  
 A>Title: Immunobistochemical and biochemical analyses of 20000-25000-year-old fossil  
 A.Reference number: S63514; MUID:96096730; PMID:8529631  
 A.Accession: S63514  
 A.Molecule type: protein  
 A.Residues: 243-261; 575-590; 756-763, 'X', '765-779 <PRA>  
 R.Tiller, G.E.; Weis, M.A.; Polunbo, P.A.; Gruber, H.E.; Rimoin, D.L.; Cohn, D.H.; Ey  
 Am. J. Hum. Genet. 56, 388-395, 1995  
 A>Title: An RNA-splicing mutation (G-51YS20) in the type II collagen gene (COL2A1) in  
 A.Reference number: I38867; MUID:95150028; PMID:7847372  
 A.Accession: I38867  
 A.Status: preliminary; translated from GB/EMBL/DBJ  
 A.Molecule type: DNA  
 A.Residues: 440, 'G', '442-456, 'E', '458-480, 'P', '482-509 <TILL>  
 A.Cross-references: EMBL:U05195; NID:9557053; PIDN:AMB60370.1; PID:9557054  
 R.Ramirez, F.  
 Submitted to the EMBL Data Library, December 1988  
 A.Reference number: S04892  
 A.Accession: S04892  
 A.Molecule type: mRNA  
 A.Residues: 501-676, 'A', '678-783, 'A', '785-831, 'PA', '834, 'F', '836-1214 <RAM>  
 A.Cross-references: EMBL:X13783; NID:930037; PIDN:CAA32030.1; PID:93003050  
 R.Vikkula, M.; Peltonen, L.  
 FEBS Lett. 250, 171-174, 1989  
 A>Title: Structural analyses of the polymorphic area in type II collagen gene.  
 A.Reference number: S05000; MUID:89325561; PMID:2753125  
 A.Accession: S05000  
 A.Molecule type: DNA  
 A.Residues: 630-640, 'A', '642-785 <VIK2>  
 A.Cross-references: EMBL:X16158; NID:929951; PIDN:CAA34278.1; PID:91335018; PIDN:CAA3  
 PIDN:CAA34283.1; PID:91335023; PIDN:CAA34284.1; PID:91335024  
 R.Bogert, R.; Tiller, G.E.; Weis, M.A.; Gruber, H.E.; Rimoin, D.L.; Cohn, D.H.; Eyre  
 J. Biol. Chem. 267, 22522-22526, 1992  
 A>Title: An amino acid substitution (G1985->S1U) in the collagen alpha 1(II) chain  
 A.Reference number: A44309; MUID:93054548; PMID:1429602  
 A.Accession: A44309  
 A.Status: nucleic acid sequence not shown; not compared with conceptual translation  
 A.Molecule type: DNA  
 A.Residues: 752-831, 'PA', '834, 'F', '836-1005, 'K', '1007-1036, 'Q', '1038-1052, 'E', '1054-1068, '  
 A.Cross-references: GB:I00977; NID:9180812; PIDN:AB23914.1; PID:9258774  
 A>Note: sequence extracted from NCBI backbone (NCBI:11773); parts of this sequence  
 A>Note: this translation is not annotated and this publication is not cited in Genban  
 A>Note: mutant sequence associated with perinatal lethal hypochondrogenesis  
 R.Tiller, G.E.; Rimoin, D.L.; Murray, L.W.; Cohn, D.H.  
 Proc. Natl. Acad. Sci. U.S.A. 87, 3869-3893, 1990  
 A>Title: Tandem duplication within a type II collagen gene (COL2A1) exon in an indivi  
 A.Reference number: S16502; MUID:90251662; PMID:2339128  
 A.Accession: S16502  
 A.Molecule type: DNA  
 A.Residues: 1164-1184, 'GPSGKGANGIPGT', '1185-1199 <TIL2>  
 A.Cross-references: EMBL:M71216; NID:9180808; PIDN:AAA52037.1; PID:9180809  
 A>Note: mutant sequence from a patient with spondyloepiphyseal dysplasia  
 R.Head, K.S.E.; Stoker, N.G.; Griffin, J.R.; Grosvear, F.G.; Solomon, E.  
 Proc. Natl. Acad. Sci. U.S.A. 82, 2555-2559, 1985  
 A>Title: Identification and characterization of the human type II collagen gene (COL2A1)  
 A.Reference number: A02858; MUID:85190534; PMID:3857598  
 A.Accession: A02858  
 A.Molecule type: DNA  
 A.Residues: 1032-1056, 'N', '1058-1068, 'T', '1070-1487 <CHE>  
 A.Cross-references: GB:J00116; NID:9180395; PIDN:AAA51997.1; PID:9180396  
 R.Elima, K.; Vuorio, T.; Vuorio, E.  
 Nucleic Acids Res. 15, 9499-9504, 1987

A:Title: Determination of the single polyadenylation site of the human pro-alpha-1(II) c  
A:Reference number: A27280; MUID:88067771; PMID:2825137  
A:Accession: A27280  
A:Molecule type: DNA; mRNA  
A:Residues: 1175-1487 <ELI>  
A:Cross-references: EMBL:X06268; NID:930096; PIDN:CAA2604.1; PID:930097  
A:Experimental source: fetal epiphyseal cartilage  
R:van der Rest, M.; Rosenberg, L.C.; Olsen, B.R.; Poole, A.R.  
Biochem. J. 237, 923-925, 1986  
A:Title: Chondrocalcin is identical with the C-propeptide of type II procollagen.  
A:Reference number: A57033; MUID:87099927; PMID:3800925  
A:Accession: A57033  
A:Molecule type: protein  
A:Residues: 'XE',1244-1246,'N',1248,'X',1250-1265;1295-1305;1395-1408 <VAN>  
A:Note: chondrocalcin identified as released collagen 1(II) chain carboxyl-terminal pro  
R:Strom, C.M.; Upholt, W.B.  
Nucleic Acids Res. 12, 1025-1038, 1984  
A:Title: Isolation and characterization of genomic clones corresponding to the human tyf  
A:Reference number: A21733; MUID:84118798; PMID:6320112  
A:Accession: A21733  
A:Molecule type: DNA  
A:Residues: 1245-1295 <STR1>  
A:Cross-references: EMBL:X00339; EMBL:X00298; NID:9394699; PIDN:CAA25092.1; PID:94378975  
A:Accession: B21733  
A:Molecule type: DNA  
A:Residues: 894-909, 'PE' <STR2>  
A:Cross-references: GB:K01785; NID:930035; PIDN:CAA25082.1; PID:91335032  
R:Nunez, A.M.; Francomano, C.; Young, M.F.; Martin, G.R.; Yamada, Y.  
Biochemistry 24, 6343-6348, 1985  
A:Title: Isolation and partial characterization of genomic clones coding for a human pro  
gene.  
A:Reference number: A24561; MUID:86104139; PMID:3002437  
A:Accession: A24561  
A:Molecule type: DNA  
A:Residues: 1296-1358 <NUN2>  
A:Cross-references: GB:M12048; NID:9180017  
A:Note: This translation is not annotated in GenBank entry HM0CCT2A, release 111.0  
R:Sangiorgi, F.O.; Benson-Chanda, V.; de Wet, W.J.; Sobel, M.E.; Tsipouras, P.; Ramirez,  
Nucleic Acids Res. 13, 2207-2225, 1985  
A:Title: Isolation and partial characterization of the entire human pro alpha 1(II) coll  
A:Reference number: I37249; MUID:85215609; PMID:2987845  
A:Accession: S59491  
A:Molecule type: DNA  
A:Residues: 7-28; 'R',99-114;541-578;786-802;1055-1056,'N',1058-1068,'T',1070-1109;1200-1  
A:Accession: I84453  
A:Molecule type: DNA  
A:Status: translated from GB/EMBL/DBJ  
A:Residues: 7-28 <SAN2>  
A:Cross-references: GB:M23759; NID:9180845; EMBL:X03320; GB:M24938; NID:930104  
A:Note: the GenBank PID is based on an incorrect reading frame  
A:Accession: I37250  
A:Status: translated from GB/EMBL/DBJ  
A:Molecule type: DNA  
A:Residues: 541-560 <SAN3>  
A:Cross-references: EMBL:X02378; GB:M23870; NID:930107; PIDN:CAA26227.1; PID:929621  
A:Accession: I37251

Query Match 88.8%; Score 71; DB 1; Length 1487;  
Best Local Similarity 80.0%; Pred. No. 0.0038;  
Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 GTPGPGIAGGRCV 15  
I |||||:||||:|  
Db 969 GPGPGIAGGRCV 983

RESULT 8  
B41182  
collagen alpha 1(II) chain precursor (long splice form) - mouse  
C:Species: Mus musculus (house mouse)  
C:Date: 28-May-1992 #sequence\_revision 28-May-1992 #text\_change 16-Jul-1999  
C:Accession: B41182

R:Meisner, M.; Toman, D.; de Crombrughe, B.; Viorio, E.  
J. Biol. Chem. 266, 16862-16869, 1991  
A:Title: Mouse type II collagen gene. Complete nucleotide sequence, exon structure, a  
A:Reference number: A41182; MUID:91358489; PMID:1885613  
A:Accession: B41182  
A:Status: preliminary; not compared with conceptual translation  
A:Molecule type: DNA  
A:Residues: 1-1487 <MET>  
A:Cross-references: GB:M65161  
C:Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homolo  
C:Keywords: alternative splicing; coiled coll; extracellular matrix; glycoprotein; tr  
F:33-91/Domain: von Willibrand factor type C repeat homology <VWC>  
F:1259-1487/Domain: fibrillar collagen carboxyl-terminal homology <FCC>

Query Match 88.8%; Score 71; DB 2; Length 1487;  
Best Local Similarity 80.0%; Pred. No. 0.0038;  
Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 GTPGPGIAGGRCV 15  
I |||||:||||:|  
Db 969 GPGPGIAGGRCV 983

RESULT 9  
B40333  
collagen alpha 1(II) chain precursor - African clawed frog  
C:Species: Xenopus laevis (African clawed frog)  
C:Date: 10-Sep-1999 #sequence\_revision 10-Sep-1999 #text\_change 10-Sep-1999  
A:Accession: B40333  
R:Su, M.W.; Suzuki, H.R.; Bleker, J.T.; Solursh, M.; Ramirez, F.  
J. Cell Biol. 115, 565-575, 1991  
A:Title: Expression of two nonallelic type II procollagen genes during Xenopus laevis  
A:Reference number: A40333; MUID:92011898; PMID:1918153  
A:Accession: B40333  
A:Status: preliminary  
A:Molecule type: mRNA  
A:Residues: 1-1486 <SUA>  
A:Cross-references: GB:M63595  
C:Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homolo  
C:Keywords: coiled coll; extracellular matrix; glycoprotein; trimer; triple helix  
F:337-96/Domain: von Willibrand factor type C repeat homology <VWC>  
F:1258-1486/Domain: fibrillar collagen carboxyl-terminal homology <FCC>

Query Match 85.0%; Score 68; DB 1; Length 1486;  
Best Local Similarity 73.3%; Pred. No. 0.011;  
Matches 11; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

OY 1 GTPGPGIAGGRCV 15  
I |||||:||||:|  
Db 971 GPGPGIAGGRCV 985

RESULT 10  
S16366  
collagen alpha 2(IV) chain precursor - pig roundworm  
C:Species: Ascaris suum (pig roundworm)  
C:Date: 04-Dec-1992 #sequence\_revision 04-Dec-1992 #text\_change 13-Aug-1999  
A:Accession: S16366  
R:Petit, J.; Kingston, I.B.  
J. Biol. Chem. 266, 16149-16156, 1991  
A:Title: The complete primary structure of a nematode alpha-2(IV) collagen and the pa  
A:Reference number: S16366; MUID:91340768; PMID:1714907  
A:Accession: S16366  
A:Molecule type: mRNA  
A:Residues: 1-1763 <JBI>  
A:Cross-references: GB:M67507; NID:9159648; PIDN:AAA18014.1; PID:9159649  
C:Genetics:  
A:introns: 229/3; 266/3; 305/3; 360/3; 424/1; 489/1; 548/1; 656/3; 790/1; 891/1; 963/3  
C:Superfamily: collagen alpha 1(IV) chain  
C:Keywords: alternative splicing; basement membrane; cell binding; coiled coll; disul  
F:27-1763/Product: collagen alpha 2(IV) chain #status predicted <MAT>  
F:27-42/Domain: non-collagenous NHI #status predicted <NHI>

F:43-1529/Domain: collagenous #status predicted <COL>  
 F:197-199/Region: cell attachment (R-G-D) motif  
 F:1530-1763/Domain: carboxyl-terminal nonhelical, NC1 #status predicted <NC1>  
 F:1530-1638/Domain: repeat NC1 #status predicted <NC1>  
 F:1639-1763/Domain: repeat NC1 #status predicted <NC1>  
 F:31,34,39,41,536,539/Disulfide bonds: Interchain #status predicted  
 F:156/Binding site: carbohydrate (Asn) (covalent) #status predicted  
 F:1593-1599,1702-1709/Disulfide bonds: #status predicted

Query Match 77.5%; Score 62; DB 2; Length 1763;  
 Best Local Similarity 66.7%; Pred. No. 0.12;  
 Matches 10; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Oy 1 GTPGPGIAGRGV 15  
 Db 82 GTPGPGIKGRGII 96

## RESULT 11

S32436  
 collagen alpha 2(IX) chain - human (fragment)  
 C:Species: Homo sapiens (man)  
 C>Date: 30-Sep-1993 #sequence\_revision 30-Sep-1993 #text\_change 08-May-1998  
 C:Accession: S32436; S34487; S64673  
 R:Peraelae, M.; Haeninen, M.; Haestbacka, J.; Elima, K.; Vuorio, E.  
 FEBS Lett. 319, 177-180, 1993  
 A>Title: Molecular cloning of the human alpha-2(IX) collagen cDNA and assignment of the  
 A:Reference number: S32436; MUID:93202262; PMID:8454052  
 A:Accession: S32436  
 A:Molecule type: mRNA  
 A:Residues: 1-618 <PER1>  
 A:Cross-references: EMBL:M95610; NID:91054872  
 R:Peraelae, M.; Haeninen, M.; Haestbacka, J.; Vuorio, E.  
 submitted to the EMBL Data Library, March 1993  
 A:Description: Molecular cloning of the human alpha-2 (IX) collagen cDNA and assignment  
 A:Reference number: S34487  
 A:Accession: S34487  
 A:Molecule type: mRNA  
 A:Residues: 1-26, 'QT', 29, 'S', 31-32, 'LM', 35-561, 'L', 563-578, 'P', 580-618 <PER2>  
 A:Cross-references: EMBL:M95610; NID:91054872  
 R:Diab, M.; Wu, J.J.; Eyre, D.R.  
 Biochem. J. 314, 327-332, 1996  
 A>Title: Collagen type IX from human cartilage: a structural profile of intermolecular c  
 A:Reference number: S64673; MUID:96195147; PMID:8660302  
 A:Accession: S64673  
 A:Molecule type: protein  
 A:Residues: 123-133, 'P', 135-137 <DIA>  
 C:Comment: Prolines and lysines at the third position of the tripeptide repeating unit o  
 ed and subsequently O-glycosylated.  
 C:Genetics:

A:Gene: GDB:COL9A2  
 A:Cross-references: GDB:138310; OMIM:120260  
 A:Map position: 1p33-1p32.2  
 C:Complex: type IX collagen may be a heterotrimer of one alpha I(IX) chain, one alpha 2  
 C:Function:  
 A:Description: structural component of extracellular fibrous polymer associated with tyf  
 C:Superfamily: unassigned collagens  
 C:Keywords: chondroitin sulfate proteoglycan; coiled coil; extracellular matrix; glycop  
 F:1-114/Domain: collagenous COL3 (fragment) #status predicted <COL3>  
 F:115-131/Domain: non-collagenous NC3 #status predicted <NC3>  
 F:133-470/Domain: collagenous COL2 #status predicted <COL2>  
 F:471-500/Domain: non-collagenous NC2 #status predicted <NC2>  
 F:501-615/Domain: collagenous COL1 #status predicted <COL1>  
 F:616-618/Domain: non-collagenous NC1 (fragment) #status predicted <NC1>  
 F:120/Binding site: chondroitin sulfate (Ser) (covalent) #status predicted

Query Match 73.8%; Score 59; DB 2; Length 618;  
 Best Local Similarity 71.4%; Pred. No. 0.13;  
 Matches 10; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Oy 1 GTPGPGIAGRGV 14  
 Db 444 GTPGPGIAGRGV 457

## RESULT 12

A32249  
 collagen - sea urchin (Paracentrotus lividus) (fragment)  
 C:Species: Paracentrotus lividus (common urchin)  
 C>Date: 17-Aug-1989 #sequence\_revision 17-Aug-1989 #text\_change 19-Jan-1996  
 C:Accession: A32249  
 R:Saltia, B.; Butlice, G.; Gambino, R.  
 Biochem. Biophys. Res. Commun. 158, 633-639, 1989  
 A>Title: Isolation of a putative collagen-like gene from the sea urchin Paracentrotus  
 A:Reference number: A32249; MUID:89149773; PMID:2537631  
 A:Accession: A32249  
 A:Status: preliminary; not compared with conceptual translation

A:Molecule type: DNA  
 A:Residues: 1-290 <SAI>  
 C:Superfamily: collagen alpha 2(I) chain; fibrillar collagen carboxyl-terminal homolo  
 C:Keywords: coiled coil; extracellular matrix; glycoprotein; trimer; triple helix

Query Match 72.5%; Score 58; DB 2; Length 290;  
 Best Local Similarity 76.9%; Pred. No. 0.09;  
 Matches 10; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Oy 1 GTPGPGIAGRG 13  
 Db 268 GTPGPGIAGRG 280

## RESULT 13

CGH02V  
 collagen alpha 2(V) chain precursor - human  
 C:Species: Homo sapiens (man)  
 C>Date: 31-Jul-1988 #sequence\_revision 28-Jul-1995 #text\_change 31-Dec-2000  
 C:Accession: A31427; A54555; S43643; A25874; I55239; I59025; A25374; A30017  
 R:Woodbury, D.; Benson-Chanda, V.; Ramirez, F.  
 J. Biol. Chem. 264, 2735-2738, 1989  
 A>Title: Amino-terminal propeptide of human pro-alpha2(V) collagen conforms to the st  
 A:Reference number: A31427; MUID:89123368; PMID:2914927  
 A:Accession: A31427  
 A:Molecule type: mRNA  
 A:Residues: 1-463 <MOO>  
 A:Cross-references: GB:J04478; NID:9179697; PIDN:AAA51859.1; PID:9179698  
 A:Experimental source: placenta  
 R:Greenspan, D.S.; Lee, S.T.; Lee, B.S.; Hoffman, G.G.  
 Gene Expr. 1, 29-39, 1991

A>Title: Homology between alpha2(V) and alpha1(III) collagen promoters and evidence f  
 A:Reference number: A54555; MUID:92314691; PMID:1820205  
 A:Accession: A54555  
 A:Molecule type: DNA  
 A:Residues: 1-32 <GRB>  
 A:Cross-references: GB:M58529; NID:9180834; PIDN:AA041699.1; PID:9553235  
 R:Moradi-Ameli, M.; Rousseau, J.C.; Klemen, J.P.; Champilaud, M.F.; Boutillon, M.M.;  
 Eur. J. Biochem. 221, 987-995, 1994  
 A>Title: Diversity in the processing events at the N-terminus of type-V collagen.  
 A:Reference number: S43642; MUID:94237164; PMID:8181482  
 A:Accession: S43642  
 A:Molecule type: protein  
 A:Residues: 288-291, 'P', 293-294, 'X', 296-297, 606, 'X', 608-617 <MOR>  
 R:Well, D.; Bernard, M.; Garcano, S.; Ramirez, F.  
 Nucleic Acids Res. 15, 181-198, 1987  
 A>Title: The pro alpha 2(V) collagen gene is evolutionarily related to the major fibr  
 A:Reference number: A25874; MUID:87146331; PMID:3025669  
 A:Accession: A25874  
 A:Molecule type: mRNA; DNA  
 A:Residues: 398-1496 <MEI>  
 A:Cross-references: GB:X04758; NID:929588; PIDN:CAA28454.1; PID:91340175  
 A:Experimental source: rhabdomyosarcoma cell line  
 R:Myers, J.C.; Loidl, H.R.; Stollé, C.A.; Seyer, J.M.  
 J. Biol. Chem. 260, 5533-5541, 1985

A>Title: Partial covalent structure of the human alpha 2 type V collagen chain.  
 A:Reference number: I55239; MUID:85182703; PMID:2985598  
 A:Accession: I55239  
 A:Status: translated from GB/EMBL/DDBT



A: Molecule type: mRNA  
 A: Residues: 1002-1226 <RES>  
 A: Cross-references: GB:M0956; NID:g180427; PIDN:AAA52007.1; PID:g180428  
 A: Note: part of this sequence were determined by protein sequencing  
 R: Emanuel, B.S.; Cammizaro, L.A.; Seyer, J.M.; Myers, J.C.  
 Proc. Natl. Acad. Sci. U.S.A. 82, 3385-3389, 1985  
 A: Title: Human alpha 1(III) and alpha 2(V) procollagen genes are located on the long arm  
 A: Reference number: 159025; MUID:85216505; PMID:3858826  
 A: Accession: 159025  
 A: Status: translated from GB/EMBL/DBDUT  
 A: Molecule type: mRNA  
 A: Residues: 1003-1034 <RES>  
 A: Cross-references: GB:M1135; NID:g179693; PIDN:AAA51857.1; PID:g179694  
 A: Note: part of this sequence were determined by protein sequencing  
 R: Myers, J.C.; Ioldi, H.R.; Seyer, J.M.; Dion, A.S.  
 J. Biol. Chem. 260, 11216-11222, 1985  
 A: Title: Complete primary structure of the human alpha-2 type V procollagen COOH-terminus  
 A: Reference number: A25374; MUID:85289337; PMID:2411731  
 A: Accession: A25374  
 A: Molecule type: mRNA  
 A: Residues: 1227-1417, 'T', 1419-1437, 'S', 1439-1496 <MYE>  
 A: Cross-references: GB:M1178; NID:g180912; PIDN:AAA52058.1; PID:g180913  
 A: Experimental source: normal fibroblasts  
 R: Tsipouras, P.; Schwartz, R.C.; Liddell, A.C.; Salkeid, C.S.; Weil, D.; Ramirez, F.  
 Genomics 3, 275-277, 1988  
 A: Title: Genetic distance of two fibrillar collagen loci, COL3A1 and COL5A2, located on  
 A: Reference number: A30017; MUID:89138450; PMID:3224983  
 A: Accession: A30017  
 A: Molecule type: DNA  
 A: Residues: 1449-1463, 'E', 1465-1495, 'A' <TST>  
 A: Cross-references: GB:U03051; NID:g179695; PIDN:AAA51858.1; PID:g179696  
 A: Note: the authors translated the codon GAA for residue 1460 as Gln, and GAG for residue  
 C: Comment: Prolines and lysines at the third position of the tripeptide repeating unit  
 are 5-hydroxylated and subsequently O-glycosylated.  
 C: Comment: The amino-terminal propeptide domain appears not to be completely cleaved.  
 C: Genetics:  
 A: Gene: GDB:COL5A2  
 A: Cross-references: GDB:119064; OMIM:120190  
 A: Map position: 2q31-2q31  
 A: Introns: 33/1; 812/3; 848/3; 884/3; 902/3; 922/3; 974/3; 1046/3; 1064/3; 1448/3  
 C: Complex: type V collagen may be a homotrimer of alpha 1(V) chains (see PIR:CGH1V), a  
 alpha 2(V) chain and one alpha 3(V) chain, initially linked by disulfide bonds among the  
 length, is formed with desmosine cross-links made from lysine and allysine residues  
 C: Function:  
 A: Description: structural component of extracellular fibrous polymer associated with cell  
 A: Note: may play a role in controlling the lateral growth of collagen I fibrils  
 C: Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homology;  
 C: Keywords: coiled coil; extracellular matrix; glycoprotein; hydroxylysine; hydroxyproline  
 F: 1-26/Domain: signal sequence #status predicted <SIG>  
 F: 27-1350/Product: collagen alpha 2(V) chain #status predicted <MAT>  
 F: 27-108/Domain: amino-terminal propeptide (uncleaved) #status predicted <NPP>  
 F: 27-108/Region: nonhelical  
 F: 40-99/Domain: von Willebrand factor type C repeat homology <WVC>  
 F: 109-186/Region: helical  
 F: 187-208/Region: nonhelical  
 F: 209-1225/Region: helical  
 F: 503-505/Region: cell attachment (R-G-D) motif  
 F: 941-943/Region: cell attachment (R-G-D) motif  
 F: 1064-1066/Region: cell attachment (R-G-D) motif  
 F: 1067-1068/Region: cell attachment (R-G-D) motif  
 F: 1097-1099/Region: cell attachment (R-G-D) motif  
 F: 1124-1126/Region: cell attachment (R-G-D) motif  
 F: 1133-1135/Region: cell attachment (R-G-D) motif  
 F: 1225-1250/Region: carboxyl-terminal nonhelical telopeptide  
 F: 1251-1496/Domain: carboxyl-terminal propeptide #status predicted <CPP>  
 F: 1269-1496/Domain: fibrillar collagen carboxyl-terminal homology <FCC>  
 F: 27/Modified site: pyrrolidone carboxylic acid (Gln) (in mature form) #status predicted  
 F: 193-194/Cleavage site: Ala-Gln (procollagen N-endopeptidase) #status predicted  
 F: 194/Modified site: pyrrolidone carboxylic acid (Gln) (in mature form) #status predicted  
 F: 201/Modified site: allysine (lys) #status predicted  
 F: 290-293-296-608-614-1004-1007-1013-1028-1034/Modified site: 4-hydroxyproline (Pro) #st  
 F: 299-1138/Modified site: 5-hydroxylysine (lys) #status predicted  
 F: 299-1139/Binding site: carbohydrate (lys) (covalent) #status predicted

F:1025/Modified site: 5-hydroxylysine (lys) #status experimental  
 F:1250-1251/Cleavage site: Glu-Asp (procollagen C-endopeptidase) #status predicted  
 F:1259-1397/Binding site: carbohydrate (asn) (covalent) #status predicted  
 F:1293-1299-1325/Disulfide bonds: interchain #status predicted  
 F:1333-1494-1402-1447/Disulfide bonds: #status predicted

Query Match 72.5%; Score 58; DB 1; Length 1496;  
 Best Local Similarity 66.7%; Pred. No. 0.45;  
 Matches 10; Conservative 1; Mismatches 4; Indels 0; Gaps 0;

OY 1 GTPPGGAGGAGV 15  
 I I I I I I I I I I I  
 Db 978 GPPGAGTTCGRCIV 992

RESULT 14  
 149607

procollagen type V alpha 2 - mouse

C: Species: Mus musculus (house mouse)

C: Date: 02-Jul-1996 #sequence\_revision 02-Jul-1996 #text\_change 13-Aug-1999

C: Accession: 149607

R: Andrikopoulos, K.; Suzuki, H.R.; Solursh, M.; Ramirez, F.  
 Dev. Dyn. 195, 113-120, 1992

A: Title: Localization of pro-alpha 2(V) collagen transcripts in the tissues of the de

A: Reference number: 149607; MUID:93214071; PMID:1297453

A: Accession: 149607

A: Status: preliminary; translated from GB/EMBL/DBDUT

A: Molecule type: mRNA

A: Residues: 1-1497 <RES>

A: Cross-references: GB:I02918; NID:g309180; PIDN:AAA37440.1; PID:g309181

C: Genetics:

A: Gene: COL5A-2

C: Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homolo

F: 39-98/Domain: von Willebrand factor type C repeat homology <WVC>

F: 120-1497/Domain: fibrillar collagen carboxyl-terminal homology <FCC>

Query Match 72.5%; Score 58; DB 2; Length 1497;  
 Best Local Similarity 66.7%; Pred. No. 0.45;  
 Matches 10; Conservative 1; Mismatches 4; Indels 0; Gaps 0;

OY 1 GTPPGGAGGAGV 15  
 I I I I I I I I I I I  
 Db 979 GPPGAGTTCGRCIV 993

RESULT 15  
 150696

collagen alpha 1(III) chain - chicken (fragment)

C: Species: Gallus gallus (chicken)

C: Date: 13-Sep-1996 #sequence\_revision 13-Sep-1996 #text\_change 13-Aug-1999

C: Accession: 150696

R: Nab, H.D.; Niu, Z.; Adams, S.L.  
 J. Biol. Chem. 269, 16443-16448, 1994

A: Title: An alternative transcript of the chick type III collagen gene that does not

A: Reference number: A54041; MUID:94266842; PMID:8206952

A: Accession: 150696

A: Status: preliminary; translated from GB/EMBL/DBDUT

A: Molecule type: mRNA

A: Residues: 1-310 <NAB>

A: Cross-references: EMBL:U07974; NID:g520456; PIDN:AAA83409.1; PID:g537433

C: Genetics:

A: Gene: COL3A1

C: Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homolo

Query Match 71.2%; Score 57; DB 2; Length 310;  
 Best Local Similarity 69.2%; Pred. No. 0.14;  
 Matches 9; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 GTPPGGAGGAGV 13  
 I I I I I I I I I I I  
 Db 275 GPPGAGTTCGRCIV 287



Search completed: August 29, 2003, 18:27:51  
Job time : 40 secs

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GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: August 29, 2003, 17:51:44 ; Search time 23 Seconds  
(without alignments)  
30.670 Million cell updates/sec

Title: US-09-935-417-1  
Perfect score: 80  
Sequence: 1 GTPGQGIAGRGV 15

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 127863 seqs, 47026705 residues

Total number of hits satisfying chosen parameters: 127863

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : SwissProt\_41\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	80	100.0	779	CA11_BOVIN	P02453 bos taurus
2	80	100.0	1453	CA11_CHICK	P02457 gallus galli
3	80	100.0	1453	CA11_MOUSE	P11087 mus musculus
4	80	100.0	1460	CA11_CANFA	O9xs17 canis famli
5	80	100.0	1464	CA11_HUMAN	P02452 homo sapien
6	71	88.8	1418	CA12_HUMAN	P02458 homo sapien
7	71	88.8	1459	CA12_MOUSE	P28481 mus musculu
8	62	77.5	1763	CA24_ASCSU	P27393 ascaris suu
9	59	73.8	689	CA29_HUMAN	O14055 homo sapien
10	58	72.5	1362	CA21_CHICK	P02467 gallus galli
11	58	72.5	1496	CA25_HUMAN	P05997 homo sapien
12	57	71.2	636	CA13_RAT	P13941 rattus norv
13	57	71.2	1049	CA13_BOVIN	P04258 bos taurus
14	57	71.2	1262	CA13_CHICK	P12105 gallus galli
15	57	71.2	1464	CA13_MOUSE	P08121 mus musculu
16	57	71.2	1466	CA13_HUMAN	P02461 mus sapien
17	56	70.0	353	CA29_CHICK	P12108 gallus galli
18	56	70.0	526	CA21_RABIT	O28668 oryctolagus
19	56	70.0	1366	CA21_CANFA	O46392 canis famli
20	56	70.0	1366	CA21_HUMAN	P08123 homo sapien
21	56	70.0	1669	CA14_HUMAN	P02462 homo sapien
22	56	70.0	1669	CA14_MOUSE	P02463 mus musculu
23	54	67.5	747	CA12_BOVIN	P02459 bos taurus
24	53	66.2	266	YXWK_CAEEL	O21184 caenorhabdi
25	53	66.2	680	CA1A_HUMAN	O00392 homo sapien
26	53	66.2	1143	CA1I_HUMAN	O14993 homo sapien
27	53	66.2	1603	CA1F_HUMAN	O07092 homo sapien
28	53	66.2	1758	CA24_CAEEL	P17140 caenorhabdi
29	52	65.0	122	CA12_RAT	P05539 rattus norv
30	52	65.0	321	CA13_BOVIN	P42916 bos taurus
31	52	65.0	369	PSPD_BOVIN	P35246 bos taurus
32	52	65.0	371	CI46_BOVIN	O8mh29 bos taurus
33	52	65.0	520	MRCO_HUMAN	O9new3 homo sapien

34	52	65.0	921	CA19_MOUSE	Q05722 mus musculu
35	52	65.0	1364	CA21_BOVIN	P02465 bos taurus
36	52	65.0	1736	CA2B_HUMAN	P13942 homo sapien
37	52	65.0	1736	CA2B_MOUSE	O64739 mus musculu
38	52	65.0	1804	CA1B_MOUSE	O61245 mus musculu
39	52	65.0	1806	CA1B_HUMAN	P12107 mus sapien
40	51	63.7	671	CA1I_RAT	P02454 rattus norv
41	51	63.7	911	CA1B_BOVIN	O28083 bos taurus
42	51	63.7	1356	CA21_ONCMY	O93484 oncorhynch
43	51	63.7	1707	CA24_MOUSE	P08122 mus musculu
44	51	63.7	1712	CA24_HUMAN	P08572 homo sapien
45	51	63.7	1775	CA14_DROME	P08120 drosophila

## ALIGNMENTS

RESULT 1	ID	CA11_BOVIN	STANDARD:	PRT:	779 AA.
AC	P02453;				
DT	21-JUL-1986 (Rel. 01, Created)				
DT	01-FEB-1994 (Rel. 28, Last sequence update)				
DT	28-FEB-2003 (Rel. 41, Last annotation update)				
DE	Collagen alpha 1(I) chain (Fragments).				
CN	COL1A1.				
OS	Bos taurus (Bovine).				
OC	Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;				
OC	Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;				
OX	Bovidae; Bovinae; Bos.				
OX	NCBI_TaxID=9913;				
RN	[1]				
RP	SEQUENCE OF 1-19.				
RA	MEDLINE=7225334; PubMed=4115172;				
RX	Rautenberg J., Jimpl R., Furtmayr H.;				
RT	"Structural characterization of N-terminal antigenic determinants in				
RT	calf and human collagen.";				
RL	Eur. J. Biochem. 27:231-237(1972).				
RN	[2]				
RP	SEQUENCE OF 20-145.				
RX	MEDLINE=76022320; PubMed=1164916;				
RA	Fietzek P.P., Kuehn K.;				
RT	"The covalent structure of collagen: amino-acid sequence of the				
RT	cyanoen-bromide peptides alpha-1-CB2, alpha-1-CB4 and alpha-1-CB5				
RT	from calf-skin collagen.";				
RL	Eur. J. Biochem. 52:77-82(1975).				
RN	[3]				
RP	SEQUENCE OF 146-294.				
RX	MEDLINE=73049499; PubMed=4673951;				
RA	Fietzek P.P., Wendt P., Keil I., Kuehn K.;				
RT	"The covalent structure of collagen: amino acid sequence of alpha-1-				
RT	CB3 from calf skin collagen.";				
RL	FEBS Lett. 26:74-76(1972).				
RN	[4]				
RP	SEQUENCE OF 295-562.				
RX	MEDLINE=74086118; PubMed=4359390;				
RA	Fietzek P.P., Rexrodt F.W., Hopper K.E., Kuehn K.;				
RT	"The covalent structure of collagen. 2. The amino-acid sequence of				
RT	alpha-1-CB7 from calf-skin collagen.";				
RL	Eur. J. Biochem. 38:396-400(1973).				
RN	[5]				
RP	SEQUENCE OF 563-675.				
RX	MEDLINE=73042276; PubMed=4343808;				
RA	Wendt P., Mark K.V.D., Rexrodt F., Kuehn K.;				
RT	"The covalent structure of collagen. The amino-acid sequence of the				
RT	112-residues. Amino-terminal part of peptide alpha-1-CB6 from calf-				
RT	skin collagen.";				
RL	Eur. J. Biochem. 30:169-183(1972).				
RN	[6]				
RP	SEQUENCE OF 676-751.				
RX	MEDLINE=73042275; PubMed=4343807;				
RA	Fietzek P.P., Rexrodt F.W., Wendt P., Stark M., Kuehn K.;				
RT	"The covalent structure of collagen. Amino-acid sequence of peptide				

RT alpha-1-CB6-C2." ;  
 RL Eur. J. Biochem. 30:163-168(1972).  
 CC -1- FUNCTION: TYPE I COLLAGEN IS A MEMBER OF GROUP I COLLAGEN  
 CC (FIBRILLAR FORMING COLLAGEN).  
 CC -1- SUBUNIT: TRIMERS OF ONE ALPHA 2(I) AND TWO ALPHA 1(I) CHAINS.  
 CC -1- TISSUE SPECIFICITY: FORMS THE FIBRILS OF TENDON, LIGAMENTS AND  
 CC BONES. IN BONES THE FIBRILS ARE MINERALIZED WITH CALCIUM  
 CC HYDROXYAPATITE.  
 CC -1- PTM: Proline residues at the third position of the tripeptide  
 CC repeating unit (G-X-Y) are hydroxylated in some or all of the  
 CC chains. Pro-726 is the only 3-hydroxypro and the only hydroxylated  
 CC proline in position X.  
 CC -1- PTM: O-linked glycan consists of a Glc-Gal disaccharide bound to  
 CC the oxygen atom of a post-translationally added hydroxyl group.  
 CC -1- MISCELLANEOUS: THE COMPLETE CHAIN CONTAINS 1052 RESIDUES.  
 DR PIR: A91193; CGB015.  
 DR InterPro: IPR000087; Collagen.  
 DR InterPro: IPR01007; VWF\_C.  
 DR ProDom: PD000007; C1g\_helix; 1.  
 DR PROSITE: PS01208; VWF\_C\_1; PARTIAL.  
 KM Extracellular matrix; Connective tissue; Repeat: Hydroxylation;  
 FT MOD\_RES 1 1  
 FT MOD\_RES 9 9  
 FT MOD\_RES 103 103  
 FT MOD\_RES 103 103  
 FT MOD\_RES 115 115  
 FT MOD\_RES 124 124  
 FT MOD\_RES 145 146  
 FT MOD\_RES 274 274  
 FT MOD\_RES 346 346  
 FT MOD\_RES 424 424  
 FT MOD\_RES 496 496  
 FT MOD\_RES 658 658  
 FT MOD\_RES 670 670  
 FT MOD\_RES 726 726  
 SQ SEQUENCE 779 AA; 70346 MW; E554A7FE084283D1 CRC64;  
 Query Match 100.0%; Score 80; DB 1; Length 779;  
 Best Local Similarity 100.0%; Pred. NO. 0.00012;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 GTPGPGIAGGGRGV 15  
 Db 509 GTPGPGIAGGGRGV 523  
 RESULT 2  
 ID CALL\_CHICK STANDARD; PRT; 1453 AA.  
 AC P02457;  
 DT 21-JUL-1986 (Rel. 01, Created)  
 DT 01-OCT-1989 (Rel. 12, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Collagen alpha 1(I) chain precursor.  
 GN COL1A1  
 OS Gallus gallus (chicken).  
 CC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 CC Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;  
 CC Gallus.  
 OX NCBI\_TaxID=9031;  
 RN [1]  
 RP SEQUENCE OF 1-153 FROM N.A.  
 RX MEDLINE=88056316; PubMed=3678834;  
 RA Flner M.H., Boedtker H., Doty P.;  
 RT "Construction and characterization of cDNA clones encoding the 5' end  
 RT of the chicken pro alpha 1(I) collagen mRNA."  
 RL Gene 56:71-78(1987).  
 RN [2]  
 RP SEQUENCE OF 1-144 FROM N.A.  
 RX MEDLINE=88007542; PubMed=2820966;  
 RA Flner M.H., Aho S., Gerstenfeld L.C., Boedtker H., Doty P.;

RT "Unusual DNA sequences located within the promoter region and the  
 RT first intron of the chicken pro-alpha 1(I) collagen gene."  
 RL J. Biol. Chem. 262:13323-13332(1987).  
 RN [3]  
 RP SEQUENCE OF 152-1187.  
 RX MEDLINE=82231995; PubMed=7093229;  
 RA Highberger J.H., Corbett C., Dixit S.N., Yu W., Seyer J.M.,  
 RA Kang A.H., Gross J.;  
 RT "Amino acid sequence of chick skin collagen alpha 1(I)-CB8 and the  
 RT complete primary structure of the helical portion of the chick skin  
 RT collagen alpha 1(I) chain."  
 RL Biochemistry 21:2048-2055(1982).  
 RN [4]  
 RP SEQUENCE OF 1200-1205.  
 RX MEDLINE=72243016; PubMed=5047697;  
 RA Eyre D.R., Glimcher M.J.;  
 RT "Evidence for a previously undetected sequence at the carboxyterminus  
 RT of the alpha 1 chain of chicken bone collagen."  
 RL Biochem. Biophys. Res. Commun. 48:720-726(1972).  
 RN [5]  
 RP SEQUENCE OF 981-1453 FROM N.A.  
 RX MEDLINE=81160715; PubMed=6927845;  
 RA Fuller F., Boedtker H.;  
 RT "Sequence determination and analysis of the 3' region of chicken pro-  
 RT alpha 1(I) and pro-alpha 2(I) collagen messenger ribonucleic acids  
 RL including the carboxy-terminal propeptide sequences."  
 RN Biochemistry 20:996-1006(1981).  
 RN [6]  
 RP SEQUENCE OF 1311-1453 FROM N.A.  
 RX MEDLINE=80134546; PubMed=6987088;  
 RA Showalter A.M., Pesciotta D.M., Eikenberry E.F., Yamamoto T.,  
 RA Pastan I., Decrombrughe B., Fietzek P.P., Olsen B.R.;  
 RT "Nucleotide sequence of a collagen cDNA-fragment coding for the  
 RT carboxyl end of pro alpha 1(I)-chains."  
 RL FEBS Lett. 111:61-65(1980).  
 CC -1- FUNCTION: TYPE I COLLAGEN IS A MEMBER OF GROUP I COLLAGEN  
 CC (FIBRILLAR FORMING COLLAGEN).  
 CC -1- SUBUNIT: TRIMERS OF ONE ALPHA 2(I) AND TWO ALPHA 1(I) CHAINS.  
 CC -1- TISSUE SPECIFICITY: FORMS THE FIBRILS OF TENDON, LIGAMENTS AND  
 CC BONES. IN BONES THE FIBRILS ARE MINERALIZED WITH CALCIUM  
 CC HYDROXYAPATITE.  
 CC -1- PTM: Proline residues at the third position of the tripeptide  
 CC repeating unit (G-X-Y) are hydroxylated in some or all of the  
 CC chains. Pro-1153 is the only 3-hydroxypro and the only  
 CC hydroxylated proline in position X.  
 CC -1- SIMILARITY: Contains 1 VWF\_C domain.  
 CC -----  
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 CC -----  
 DR EMBL: M17839; AAA48704.1; -;  
 DR EMBL: M17838; AAA48704.1; JOINED.  
 DR EMBL: V00401; CAA23695.1; -;  
 DR EMBL: M10571; AAA48671.1; ALT-SEQ.  
 DR EMBL: M17607; AAA48672.1; -;  
 DR PIR: A27179; A27179.  
 DR PIR: I50629; I50629.  
 DR InterPro: IPR000087; Collagen.  
 DR InterPro: IPR000885; Fib\_collagen\_C.  
 DR InterPro: IPR001007; VWF\_C.  
 DR Pfam: PF01410; COLF1; 1.  
 DR Pfam: PF01391; Collagen; 18.  
 DR Pfam: PF00093; vwc; 1.  
 DR ProDom: PD000007; C1g\_helix; 2.  
 DR ProDom: PD002078; Fib\_collagen\_C; 1.  
 DR SMART: SM00038; COLF1; 1.  
 DR SMART: SM00214; VWF\_C; 1.  
 DR PROSITE: PS01208; VWF\_C\_1; 1.

DR PROSITE: PS50184; WVFC\_2; 1.  
 KW Extracellular matrix; Connective tissue; Repeat; Hydroxylation;  
 KW Glycoprotein; Collagen; Signal; Pyrrolidone carboxylic acid.  
 FT SIGNAL 1 22  
 FT PROPEP 23 151 AMINO-TERMINAL PROPEPTIDE.  
 FT CHAIN 152 1205 COLLAGEN ALPHA 1(I) CHAIN.  
 FT PROPEP 1206 1453 C-TERMINAL PROPEPTIDE.  
 FT DOMAIN 31 89 WVFC.  
 FT MOD\_RES 152 152 PYRROLIDONE CARBOXYLIC ACID.  
 FT MOD\_RES 254 254 HYDROXYLATION (POTENTIAL).  
 FT MOD\_RES 851 851 HYDROXYLATION (POTENTIAL).  
 FT MOD\_RES 1081 1081 HYDROXYLATION (POTENTIAL).  
 FT MOD\_RES 1097 1097 HYDROXYLATION (POTENTIAL).  
 FT MOD\_RES 1153 1153 HYDROXYLATION.  
 FT CONFLICT 1187 1187 F -> L (IN REF. 5).  
 FT CONFLICT 1441 1441 O -> H (IN REF. 6).  
 SQ SEQUENCE 1453 AA; 137789 MW; 3BC6152134271F4D CRC64;  
 Query Match 100.0%; Score 80; DB 1; Length 1453;  
 Best Local Similarity 100.0%; Pred. No. 0.00022;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 GTPGPGIAGGGRGV 15  
 DB 936 GTPGPGIAGGGRGV 950

RESULT 3  
 CALL\_MOUSE STANDARD; PRT: 1453 AA.  
 ID CALL\_MOUSE  
 AC P11087; O60635; (Created)  
 DT 01-JUL-1989 (Rel. 11, Last sequence update)  
 DT 01-NOV-1997 (Rel. 35, Last sequence update)  
 DT 26-FEB-2003 (Rel. 41, Last annotation update)  
 DE Collagen alpha 1(I) chain precursor.  
 GN COL1A1 OR COL1A1.  
 OS Mus musculus (Mouse).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 CC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
 OX NCBI\_TaxID=10090;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC STRAIN=FVB/N;  
 RX MEDLINE=96033240; PubMed=8535610;  
 RA Li S.W., Khillan J., Prockop D.J.;  
 RT "The complete cDNA coding sequence for the mouse pro alpha 1(I) chain  
 of type I procollagen."  
 RT Matrix Biol. 14:593-595(1995).  
 RN [2]  
 RP SEQUENCE OF 518-1128 FROM N.A.  
 RX MEDLINE=86137403; PubMed=3841523;  
 RA French B.T., Lee W.-H., Maul G.G.;  
 RT "Nucleotide sequence of a cDNA clone for mouse pro alpha 1(I)  
 collagen protein."  
 RT Gene 39:311-312(1985).  
 RN [3]  
 RP SEQUENCE OF 735-1130 FROM N.A.  
 RX MEDLINE=83141374; PubMed=6298597;  
 RA Monson J.M., Friedman J., McCarthy B.J.;  
 RT "DNA sequence analysis of a mouse pro alpha 1(I) procollagen gene:  
 evidence for a mouse B1 element within the gene."  
 RT Mol. Cell. Biol. 2:1362-1371(1982).  
 RN [4]  
 RP SEQUENCE OF 735-878 AND 1005-1058 FROM N.A.  
 RX MEDLINE=83157109; PubMed=6219867;  
 RA Monson J.M., McCarthy B.J.;  
 RT "Identification of a Balb/c mouse pro alpha 1(I) procollagen gene:  
 evidence for insertions or deletions in gene coding sequences."  
 RT DNA 1:59-69(1981).  
 RN [5]  
 RP SEQUENCE OF 1442-1453 FROM N.A.  
 RX MEDLINE=88124276; PubMed=3340560;  
 RA Mooslehner K., Harbers K.;

FT "Two mRNAs of mouse pro alpha 1(I) collagen gene differ in the size  
 RT of the 3'-untranslated region."  
 RL Nucleic Acids Res. 16:773-773(1988).  
 CC  
 CC -I- FUNCTION: TYPE I COLLAGEN IS A MEMBER OF GROUP I COLLAGEN  
 CC (FIBRILLAR FORMING COLLAGEN).  
 CC -I- SUBUNIT: TRIMERS OF ONE ALPHA 2(I) AND TWO ALPHA 1(I) CHAINS.  
 CC -I- TISSUE SPECIFICITY: FORMS THE FIBRILS OF TENDON, LIGAMENTS AND  
 CC BONES. IN BONES THE FIBRILS ARE MINERALIZED WITH CALCIUM  
 CC HYDROXAPATITE.  
 CC -I- PIV: Prolines at the third position of the tripeptide repeating  
 CC unit (G-X-Y) are hydroxylated in some or all of the chains.  
 CC -I- SIMILARITY: Contains 1 WVFC domain.  
 CC  
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 CC  
 CC EMBL, U08020; AAA88912.1; -  
 CC EMBL, X15896; CAA33904.1; -  
 CC EMBL, M14423; AAA37333.1; -  
 CC EMBL, M17491; AAA37334.1; -  
 CC EMBL, X06753; CAA29927.1; -  
 CC EMBL, X03036; AAA37332.1; -  
 CC EMBL, K03029; AAA37332.1; JOINED.  
 CC EMBL, K03030; AAA37332.1; JOINED.  
 CC EMBL, K03031; AAA37332.1; JOINED.  
 CC EMBL, K03032; AAA37332.1; JOINED.  
 CC EMBL, K03033; AAA37332.1; JOINED.  
 CC EMBL, K03034; AAA37332.1; JOINED.  
 CC EMBL, K03035; AAA37332.1; JOINED.  
 CC PIR, S57243; S21626.  
 CC MGD, MGI:88467; Colla1.  
 CC InterPro: IPR000087; Collagen.  
 CC InterPro: IPR000885; Fib collagen\_C.  
 CC InterPro: IPR001007; WVFC.  
 CC Pfam: PF01410; COLFI; 1.  
 CC ProDom: PD000007; C1g\_helix; 1.  
 CC ProDom: PD002078; Fib\_collagen\_C; 1.  
 CC SMART: SM00038; COLFI; 1.  
 CC SMART: SM00214; WVFC; 1.  
 DR PROSITE: PS01208; WVFC\_1; 1.  
 DR PROSITE: PS50184; WVFC\_2; 1.  
 KW Extracellular matrix; Connective tissue; Repeat; Hydroxylation;  
 KW Glycoprotein; Collagen; Signal.  
 FT SIGNAL 1 22  
 FT PROPEP 23 151 AMINO-TERMINAL PROPEPTIDE.  
 FT CHAIN 152 1207 COLLAGEN ALPHA 1(I) CHAIN.  
 FT PROPEP 1208 1453 CARBOXYL-TERMINAL PROPEPTIDE.  
 FT DOMAIN 29 87 WVFC.  
 FT DOMAIN 152 167 NONHELICAL REGION (N-TERMINAL).  
 FT DOMAIN 168 181 TRIPLE-HELICAL REGION.  
 FT DOMAIN 1182 1207 NONHELICAL REGION (C-TERMINAL).  
 FT CARBOHD 56 56 N-LINKED (GLCNAC. . .) (POTENTIAL).  
 FT CARBOHD 1354 1354 N-LINKED (GLCNAC. . .) (POTENTIAL).  
 FT SITE 734 736 CELL ATTACHMENT SITE (POTENTIAL).  
 FT SITE 1082 1084 CELL ATTACHMENT SITE (POTENTIAL).  
 FT CONFLICT 1450 1450 A -> V (IN REF. 5).  
 SQ SEQUENCE 1453 AA; 137944 MW; 3B802E535DF81608 CRC64;  
 Query Match 100.0%; Score 80; DB 1; Length 1453;  
 Best Local Similarity 100.0%; Pred. No. 0.00022;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 GTPGPGIAGGGRGV 15  
 DB 936 GTPGPGIAGGGRGV 950

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RESULT 4
CALL_CANFA STANDARD: PRT: 1460 AA.
ID CALL_CANFA
AC 09XSJ7:
DT 30-MAY-2000 (Rel. 39, Created)
DT 30-MAY-2000 (Rel. 39, Last sequence update)
DT 30-MAY-2000 (Rel. 39, Last annotation update)
DE Collagen alpha 1(I) chain precursor.
GN COL1A1.
OS Canis familiaris (Dog).
OC Eukaryota; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Canis.
OX NCBI_TaxID=9615;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Skin;
RA Campbell B.G., Wootton J.A.M., McLeod J.N., Minor R.R.;
RL "Sequence of normal canine COL1A1 cDNA.";
RL Submitted (MAY-1999) to the EMBL/Genbank/DBJ databases.
CC -1- FUNCTION: TYPE I COLLAGEN IS A MEMBER OF GROUP I COLLAGEN
CC (FIBRILLAR FORMING COLLAGEN).
CC -1- SUBUNIT: TRIMERS OF ONE ALPHA 2(I) AND TWO ALPHA 1(I) CHAINS.
CC -1- PM: Prolines at the third position of the tripeptide repeating
CC unit (G-X-Y) are hydroxylated in some or all of the chains.
CC -1- SIMILARITY: Contains 1 WFPC domain.
CC -----
CC This SWISS-PROT entry is copyright. It is produced through a collaboration
CC between the Swiss Institute of Bioinformatics and the EMBL Outstation -
CC the European Bioinformatics Institute. There are no restrictions on its
CC use by non-profit institutions as long as its content is in no way
CC modified and this statement is not removed. Usage by and for commercial
CC entities requires a license agreement (See http://www.isb-sib.ch/announce/
CC or send an email to license@isb-sib.ch).
CC -----
DR EMBL: AF153062; AAD4619.1;
DR InterPro: IPR000087; Collagen.
DR InterPro: IPR000885; Fib_collagen_C.
DR InterPro: IPR001007; WF_C.
DR Pfam: PF01410; COLFI; 1.
DR Pfam: PF01391; Collagen; 18.
DR ProDom: PD000007; Ctg_helix; 2.
DR ProDom: PD002078; Fib_collagen_C; 1.
DR SMART: SM00038; COLFI; 1.
DR SMART: SM00214; WVC; 1.
DR PROSITE: PS01208; WVC_1; 1.
DR PROSITE: PS50184; WVC_2; 1.
KW Extracellular matrix; Connective tissue; Repeat; Hydroxylation;
KW Glycoprotein; Collagen; Signal.
FT SIGNAL 1 22
FT PROPEP 23 157 BY SIMILARITY.
FT CHAIN 158 1214 AMINO-TERMINAL PROPEPTIDE.
FT PROPEP 1215 1460 COLLAGEN ALPHA 1(I) CHAIN.
FT DOMAIN 34 92 CARBOXYL-TERMINAL PROPEPTIDE.
FT DOMAIN 158 174 WFPC.
FT DOMAIN 175 1186 NONHELICAL REGION (N-TERMINAL).
FT DOMAIN 1189 1214 TRIPLE-HELICAL REGION.
FT SITE 741 743 NONHELICAL REGION (C-TERMINAL).
FT SITE 1089 1091 CELL ATTACHMENT SITE (POTENTIAL).
FT SITE 1361 1361 CELL ATTACHMENT SITE (POTENTIAL).
FT CARBOHYD N-LINKED (GLCNAC) (POTENTIAL).
SQ SEQUENCE 1460 AA: 138762 MW: 585674D2570697 CRC64;

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Query Match 100.0%; Score 80; DB 1; Length 1460;
Best Local Similarity 100.0%; Pred. No. 0.00023;
Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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OY 1 GTRGPQGIAGRGVY 15
DB 943 GTRGPQGIAGRGVY 957

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RESULT 5
CALL_HUMAN STANDARD: PRT: 1464 AA.
ID CALL_HUMAN

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AC P02452; Q14037; Q15176;
DT 21-JUL-1986 (Rel. 01, Created)
DT 01-MAR-1989 (Rel. 10, Last sequence update)
DT 28-FEB-2003 (Rel. 41, Last annotation update)
DE Collagen alpha 1(I) chain precursor.
GN COL1A1.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE OF 1-472 FROM N.A.
RX MEDLINE=89025644; PubMed=3178743;
RA Tromp G., Kuivaniemi H., Stacey A., Shikata H., Baldwin C.T.,
RA Jaenisch R., Prockup D.J.;
RL "Structure of a full-length cDNA clone for the prepro alpha 1(I)
RL chain of human type I procollagen.";
RL Biochem. J. 253:919-922(1988).
RN [2]
RP SEQUENCE OF 1-181 FROM N.A.
RX MEDLINE=84270697; PubMed=6462220;
RA Chu M.-L., de Wet W.J., Bernard M.P., Ding J.-F., Morabito M.,
RA Myers J., Williams C., Ramirez F.;
RL "Human pro alpha 1(I) collagen gene structure reveals evolutionary
RL conservation of a pattern of introns and exons.";
RL Nature 310:337-340(1984).
RN [3]
RP SEQUENCE OF 162-301.
RC TISSUE=Skin;
RX MEDLINE=71038625; PubMed=5529814;
RA Click E.M., Bornstein P.;
RL "Isolation and characterization of the cyanogen bromide peptides from
RL the alpha 1 and alpha 2 chains of human skin collagen.";
RL Biochemistry 9:4699-4706(1970).
RN [4]
RP SEQUENCE OF 263-268.
RC TISSUE=Skin;
RX MEDLINE=71001508; PubMed=4319110;
RA Morgan P.H., Jacobs H.G., Segrest J.P., Cunningham L.W.;
RL "A comparative study of glycopeptides derived from selected
RL vertebrate collagens. A possible role of the carbohydrate in fibril
RL formation.";
RL J. Biol. Chem. 245:5042-5048(1970).
RN [5]
RP SEQUENCE OF 425-1464 FROM N.A.
RX MEDLINE=84080385; PubMed=6689127;
RA Bernard M.P., Chu M.-L., Myers J.C., Ramirez F., Eikenberry E.F.,
RA Prockup D.J.;
RL "Nucleotide sequences of complementary deoxyribonucleic acids for the
RL pro alpha 1 chain of human type I procollagen. Statistical evaluation
RL of structures that are conserved during evolution.";
RL Biochemistry 22:5213-5223(1983).
RN [6]
RP SEQUENCE OF 1229-1454 FROM N.A.
RC TISSUE=Bone;
RX MEDLINE=88124208; PubMed=3340531;
RA Maekelae J.K., Raassina M., Virta A., Vuorio E.;
RL "Human pro alpha 1(I) collagen: cDNA sequence for the C-propeptide
RL domain.";
RL Nucleic Acids Res. 16:349-349(1988).
RN [7]
RP SEQUENCE OF 1-34 FROM N.A.
RX MEDLINE=88097389; PubMed=3480516;
RA Bornstein P., McKay J., Morishima J.K., Devarayalu S., Gellinas R.E.;
RL "Regulatory elements in the first intron contribute to
RL transcriptional control of the human alpha 1(I) collagen gene.";
RL Proc. Natl. Acad. Sci. U.S.A. 84:8869-8873(1987).
RN [8]
RP SEQUENCE OF 1-34 FROM N.A.
RX MEDLINE=85130970; PubMed=2857713;
RA Chu M.-L., de Wet W.J., Bernard M.P., Ramirez F.;
RL "Fine structural analysis of the human pro-alpha 1 (I) collagen gene.
RL Promoter structure, Alu repeats, and polymorphic transcripts.";

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RL J. Biol. Chem. 260:2315-2320(1985).  
 RN [9]  
 RP SEQUENCE OF 1-44 FROM N.A.  
 RX MEDLINE-88033098; PubMed-2822714;  
 RA Rossouw C.M.S., Vergeer W.P., du Plooy S.J., Bernard M.P., Ramirez F.,  
 de Wet W.J.;  
 RT "DNA sequences in the first intron of the human pro-alpha 1(I)  
 collagen gene enhance transcription.";  
 RL J. Biol. Chem. 262:15151-15157(1987).  
 RN [10]  
 RP REVIEW ON VARIANTS.  
 RX MEDLINE-91184577; PubMed-2010058;  
 RA Kulvanliemi H., Tromp G., Prockop D.J.;  
 RT "Mutations in collagen genes: causes of rare and some common diseases  
 in humans.";  
 RL FASEB J. 5:2052-2060(1991).  
 RN [11]  
 RP REVIEW ON VARIANTS.  
 RX MEDLINE-97255959; PubMed-9101290;  
 RA Kulvanliemi H., Tromp G., Prockop D.J.;  
 RT "Mutations in fibrillar collagens (types I, II, III, and XI), fibril-  
 associated collagen (type IX), and network-forming collagen (type X)  
 cause a spectrum of diseases of bone, cartilage, and blood vessels.";  
 RL Hum. Mutat. 9:300-315(1997).  
 RN [12]  
 RP REVIEW ON VARIANTS.  
 RX MEDLINE-91374476; PubMed-1895312;  
 RA Byers P.H., Wallis G.A., Willing M.C.;  
 RT "Osteogenesis imperfecta: translation of mutation to phenotype.";  
 RL J. Med. Genet. 28:433-442(1991).  
 RN [13]  
 RP REVIEW ON VARIANTS.  
 RX MEDLINE-97169389; PubMed-9016532;  
 RA Dalglish R.;  
 RT "The human type I collagen mutation database.";  
 RL Nucleic Acids Res. 25:181-187(1997).  
 RN [14]  
 RP VARIANT OI-II CYS-1166.  
 RX MEDLINE-86287390; PubMed-3016737;  
 RA Cohn D.H., Byers P.H., Steinmann B., Gellinas R.E.;  
 RT "Lethal osteogenesis imperfecta resulting from a single nucleotide  
 change in one human pro alpha 1(I) collagen allele.";  
 RL Proc. Natl. Acad. Sci. U.S.A. 83:6045-6047(1986).  
 RN [15]  
 RP VARIANT OI-II ARG-569.  
 RX MEDLINE-87222293; PubMed-3108247;  
 RA Bateman J.F., Chan D., Walkers I.D., Rogers J.G., Cole W.G.;  
 RT "Lethal perinatal osteogenesis imperfecta due to the substitution of  
 arginine for glycine at residue 391 of the alpha 1(I) chain of type I  
 collagen.";  
 RL J. Biol. Chem. 262:7021-7027(1987).  
 RN [16]  
 RP VARIANT OI-II CYS-926.  
 RX MEDLINE-88033031; PubMed-3667599;  
 RA Vogel B.E., Minor R.R., Freund M., Prockop D.J.;  
 RT "A point mutation in a type I procollagen gene converts glycine 748  
 of the alpha 1 chain to cysteine and destabilizes the triple helix in  
 a lethal variant of osteogenesis imperfecta.";  
 RL J. Biol. Chem. 262:14737-14744(1987).  
 RN [17]  
 RP VARIANT OI-II ARG-842.  
 RX MEDLINE-86298828; PubMed-3403550;  
 RA Bateman J.F., Lamanche S.R., Dahl H.H., Chan D., Cole W.G.;  
 RT "Substitution of arginine for glycine 664 in the collagen alpha 1(I)  
 chain in lethal perinatal osteogenesis imperfecta. Demonstration of  
 the peptide defect by in vitro expression of the mutant cDNA.";  
 RL J. Biol. Chem. 263:11627-11630(1988).  
 RN [18]  
 RP VARIANT OI CYS-1195.  
 RX MEDLINE-89218628; PubMed-3244312;  
 RA Labhard M.E., Wirtz M.K., Pope F.M., Nicholls A.C., Hollister D.W.;  
 RT "A cysteine for glycine substitution at position 1017 in an alpha  
 1(I) chain of type I collagen in a patient with mild dominantly

RT inherited osteogenesis imperfecta.";  
 RL Mol. Biol. Med. 5:197-207(1988).  
 RN [19]  
 RP VARIANT OI-II VAL-434.  
 RX MEDLINE-89255493; PubMed-2470760;  
 RA Patterson E., Smiley E., Bonadio J.;  
 RT "RNA sequence analysis of a perinatal lethal osteogenesis imperfecta  
 mutation.";  
 RL J. Biol. Chem. 264:10083-10087(1989).  
 RN [20]  
 RP VARIANT OI-IV SER-1010.  
 RX MEDLINE-89308591; PubMed-2745420;  
 RA Marini J.C., Grange D.K., Gottesman G.S., Lewis M.B., Koepf D.A.;  
 RT "Osteogenesis imperfecta type IV. Detection of a point mutation in  
 one alpha 1(I) collagen allele (COL1A1) by RNA/RNA hybrid analysis.";  
 RL J. Biol. Chem. 264:11893-11900(1989).  
 RN [21]  
 RP VARIANTS OI-II ALA-1106; VAL-1151; ARG-1154 AND VAL-1184.  
 RX MEDLINE-89380165; PubMed-2777764;  
 RA Lamanche S.R., Dahl H.H., Cole W.G., Bateman J.F.;  
 RT "Characterization of point mutations in the collagen COL1A1 and  
 COL1A2 genes causing lethal perinatal osteogenesis imperfecta.";  
 RL J. Biol. Chem. 264:15809-15812(1989).  
 RN [22]  
 RP VARIANT OI SER-1022.  
 RX MEDLINE-90062068; PubMed-2511192;  
 RA Pack M., Constantinou C.D., Kalla K., Nielsen K.B., Prockop D.J.;  
 RT "Substitution of serine for alpha 1(I)-glycine 844 in a severe  
 variant of osteogenesis imperfecta minimally destabilizes the triple  
 helix of type I procollagen. The effects of glycine substitutions on  
 thermal stability are either position of amino acid specific.";  
 RL J. Biol. Chem. 264:19694-19699(1989).  
 RN [23]  
 RP VARIANT OI-II CYS-1082.  
 RX MEDLINE-89109573; PubMed-2913053;  
 RA Constantinou C.D., Nielsen K.B., Prockop D.J.;  
 RT "A lethal variant of osteogenesis imperfecta has a single base  
 mutation that substitutes cysteine for glycine 904 of the alpha 1(I)  
 chain of type I procollagen. The asymptomatic mother has an  
 unidentified mutation producing an overmodified and unstable type I  
 procollagen.";  
 RL J. Clin. Invest. 83:574-584(1989).  
 RN [24]  
 RP VARIANTS OI CYS-272; CYS-704 AND CYS-896.  
 RX MEDLINE-90009313; PubMed-2794057;  
 RA Starman B.J., Eyre D., Charbonneau H., Harrylock M., Weis M.A.,  
 RA Weiss L., Graham J.M., Byers P.H.;  
 RT "Osteogenesis imperfecta. The position of substitution for glycine by  
 cysteine in the triple helical domain of the pro alpha 1(I) chains of  
 type I collagen determines the clinical phenotype.";  
 RL J. Clin. Invest. 84:1206-1214(1989).  
 RN [25]  
 RP VARIANT OI-II CYS-422.  
 QY Query Match 100.0%; Score 80; DB 1; Length 1464;  
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 DB 947 GTPGPGIAGRGV 15  
 1 GTPGPGIAGRGV 15  
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 RESULT 6  
 CA12 HUMAN STANDARD; PRT; 1418 AA.  
 ID CA12 HUMAN  
 AC P02458;  
 DT 21-JUL-1986 (Rel. 01, Created)  
 DT 01-JAN-1990 (Rel. 13, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Collagen alpha 1(I) chain precursor (Contains: Chondrocalcin).  
 GN COL2A1.  
 OS Homo sapiens (Human).

CC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Primates; Carnivora; Hominoidea; Homo.  
 NCBI\_TaxID:9606;  
 [1]  
 RP SEQUENCE FROM N.A.  
 RX MEDLINE-90067946; PubMed-2587267;  
 RA Su M.W., Lee B., Ramirez F., Machado M., Horton W.;  
 RT "Nucleotide sequence of the full length cDNA encoding for human type  
 II procollagen.";  
 RL Nucleic Acids Res. 17:9473-9473(1989).  
 [2]  
 RP SEQUENCE OF 1-28 FROM N.A.  
 RX MEDLINE-87031574; PubMed-3021582;  
 RA Nunez A.M., Kohno K., Martin G.R., Yamada Y.;  
 RT "Promoter region of the human pro-alpha 1(II)-collagen gene.";  
 RL Gene 44:11-16(1986).  
 [3]  
 RP SEQUENCE OF 432-1145 FROM N.A.  
 RA Ramirez F.;  
 RT Submitted (DEC-1988) to the EMBL/GenBank/DBJ databases.  
 [4]  
 RP SEQUENCE OF 963-1418 FROM N.A.  
 RX MEDLINE-85190534; PubMed-3857598;  
 RA Cheah K.S.E., Stoker N.G., Griffin J.R., Grosfeld F.G., Solomon E.;  
 RT "Identification and characterization of the human type II collagen  
 gene (COL2A1).";  
 RL Proc. Natl. Acad. Sci. U.S.A. 82:2555-2559(1985).  
 [5]  
 RP SEQUENCE OF 1120-1398 FROM N.A.  
 RX MEDLINE-85306861; PubMed-3840017;  
 RA Elima K., Maekela J.K., Vuorio T., Kauppinen S., Knowles J.,  
 RT Vuorio E.;  
 RT "Construction and identification of a cDNA clone for human type II  
 procollagen mRNA.";  
 RL Biochem. J. 229:183-188(1985).  
 [6]  
 RP SEQUENCE OF 1106-1418 FROM N.A.  
 RX MEDLINE-88067771; PubMed-2825137;  
 RA Elima K., Vuorio T., Vuorio E.;  
 RT "Determination of the single polyadenylation site of the human pro  
 alpha 1(II) collagen gene.";  
 RL Nucleic Acids Res. 15:9499-9504(1987).  
 [7]  
 RP SEQUENCE OF 1227-1289 FROM N.A.  
 RX MEDLINE-86104139; PubMed-3002437;  
 RA Nunez A.M., Francosano C., Young M.F., Martin G.R., Yamada Y.;  
 RT "Isolation and partial characterization of genomic clones coding for  
 a human pro-alpha 1 (II) collagen chain and demonstration of  
 restriction fragment length polymorphism at the 3' end of the gene.";  
 RL Biochemistry 24:6343-6348(1985).  
 [8]  
 RP SEQUENCE OF 1176-1226 FROM N.A.  
 RX MEDLINE-84118798; PubMed-6320112;  
 RA Strom C.M., Upholt W.B.;  
 RT "Isolation and characterization of genomic clones corresponding to  
 the human type II procollagen gene.";  
 RL Nucleic Acids Res. 12:1025-1038(1984).  
 [9]  
 RP SEQUENCE OF 35-167 FROM N.A.  
 RX MEDLINE-89231318; PubMed-2714801;  
 RA Su M.W., Benson-Chanda V., Vissing H., Ramirez F.;  
 RT "Organization of the exons coding for pro alpha 1(II) collagen N-  
 propeptide confirms a distinct evolutionary history of this domain of  
 the fibrillar collagen genes.";  
 RL Genomics 4:438-441(1989).  
 [10]  
 RP REVIEW ON VARIANTS.  
 RX MEDLINE-91184577; PubMed-2010058;  
 RA Kuivaniemi H., Tromp G., Prockop D.J.;  
 RT "Mutations in collagen genes: causes of rare and some common diseases  
 in humans.";  
 RL FASEB J. 5:2052-2060(1991).  
 [11]

RP REVIEW ON VARIANTS.  
 RX MEDLINE-97255959; PubMed-9101290;  
 RA Kuivaniemi H., Tromp G., Prockop D.J.;  
 RT "Mutations in fibrillar collagens (types I, II, III, and XI), fibril-  
 associated collagen (type IX), and network-forming collagen (type X)  
 cause a spectrum of diseases of bone, cartilage, and blood vessels.";  
 RL Hum. Mutat. 9:300-315(1997).  
 [12]  
 RP VARIANT SER-1074.  
 RX MEDLINE-90036909; PubMed-2572591;  
 RA Vissing H., D'Alessio M., Lee B., Ramirez F., Godfrey M.,  
 RT Hollister D.W.;  
 RT "Glycine to serine substitution in the triple helical domain of pro-  
 alpha 1 (II) collagen results in a lethal perinatal form of short-  
 limbed dwarfism.";  
 RL J. Biol. Chem. 264:18265-18267(1989).  
 [13]  
 RP VARIANT SEDC 1095-GLY--TYR-1330 DEL.  
 RX MEDLINE-89266907; PubMed-2543071;  
 RA Lee B., Vissing H., Ramirez F., Rogers D., Rimoin D.;  
 RT "Identification of the molecular defect in a family with  
 spondyloepiphyseal dysplasia.";  
 RL Science 244:978-980(1989).  
 [14]  
 RP VARIANT OSTEOARTHRITIS CYS-650.  
 RX MEDLINE-90370826; PubMed-1975693;  
 RA Ala-Kokko L., Baldwin C.T., Moskowitz R.W., Prockop D.J.;  
 RT "Single base mutation in the type II procollagen gene (COL2A1) as a  
 cause of primary osteoarthritis associated with a mild  
 chondrodysplasia.";  
 RL Proc. Natl. Acad. Sci. U.S.A. 87:6565-6568(1990).  
 [15]  
 RP VARIANT OI-IV VAL-717.  
 RX MEDLINE-91291136; PubMed-2064612;  
 RA Bateman J.F., Hannagan M., Chan D., Cole W.G.;  
 RT "Characterization of a type I collagen alpha 2(I) glycine-586 to  
 valine substitution in osteogenesis imperfecta type IV. Detection of  
 the mutation and prenatal diagnosis by a chemical cleavage method.";  
 RL Biochem. J. 276:765-770(1991).  
 [16]  
 RP VARIANT OSTEOARTHRITIS CYS-650.  
 RX MEDLINE-91086471; PubMed-1985108;  
 RA Eyre D.R., Weis M.A., Moskowitz R.W.;  
 RT "Cartilage expression of a type II collagen mutation in an inherited  
 form of osteoarthritis associated with a mild chondrodysplasia.";  
 RL J. Clin. Invest. 87:357-361(1991).  
 [17]  
 RP VARIANT HYPOCHONDROGENESIS GLU-984.  
 RX MEDLINE-93054548; PubMed-1429602;  
 RA Bogaert R., Tiller G.E., Wiles M.A., Gruber H.E., Rimoin D.L.,  
 RT Cohn D.H., Eyre D.R.;  
 RT "An amino acid substitution (Gly983-->Glu) in the collagen alpha  
 1(II) chain produces hypochondrogenesis.";  
 RL J. Biol. Chem. 267:22522-22526(1992).  
 [18]  
 RP VARIANT HYPOCHONDROGENESIS SER-705.  
 RX MEDLINE-92262484; PubMed-1374906;  
 RA Horton W.A., Machado M.A., Ellard J., Campbell D., Bartley J.,  
 RA Ramirez F., Vitale E., Lee B.;  
 RT "Characterization of a type II collagen gene (COL2A1) mutation  
 identified in cultured chondrocytes from human hypochondrogenesis.";  
 RL Proc. Natl. Acad. Sci. U.S.A. 89:4583-4587(1992).  
 [19]  
 RP VARIANT WS-II ASP-198.  
 RX MEDLINE-93304428; PubMed-8317498;  
 RA Koerkoe J., Ritvanen P., Haataja L., Kaeerlahti H.,  
 RA Kuivaniemi H., Prockop D.J., Ala-Kokko L.;  
 RT "Mutation in type II procollagen (COL2A1) that substitutes aspartate  
 for glycine alpha 1-67 and that causes cataracts and retinal  
 detachment: evidence for molecular heterogeneity in the Wagner  
 syndrome and the Stickler syndrome (arthro-ophthalmopathy).";  
 RL Am. J. Hum. Genet. 53:55-61(1993).  
 [20]



RP VARIANT SEMD CYS-840.  
 RA Tyller G.E., Wells M.A., Lachman R.S., Cohn D.H., Rimoin D.L.,  
 RT "A dominant mutation in the type II collagen gene (COL2A1) produces  
 RT spondyloepiphyseal dysplasia (SEMD), Strudwick type-";  
 RL Am. J. Hum. Genet. 53:A209-A209(1993).  
 RN [21]  
 RP VARIANT OSTEOARTHRTIS CYS-650.  
 RX MEDLINE-93282819; PubMed-8507190;  
 RA Holderbaum D., Malemud C.J., Moskowitz R.W., Haq I.T.M.;  
 RT "Human cartilage from late stage familial osteoarthritis transcribes  
 RT type II collagen mRNA encoding a cysteine in position 519."  
 RL Biochem. Biophys. Res. Commun. 192:1169-1174(1993).  
 RN [22]  
 RP VARIANT SEMD ARG-285.  
 RX MEDLINE-93252400; PubMed-8486375;  
 RA Vilkula M., Rittvanleml P., Vuorio A.F., Kaitila I., Ala-Kokko L.,  
 RT Peltonen L.;  
 RT "A mutation in the amino-terminal end of the triple helix of type II  
 RT collagen causing severe osteochondrodysplasia.";  
 RL Genomics 16:282-285(1993).  
 RN [23]  
 RP VARIANT SEDC CYS-206.  
 RX MEDLINE-94063862; PubMed-8244341;  
 RA Williams C.J., Considine E.L., Knowlton R.G., Reginato A., Neumann G.,  
 RT Harrison D., Buxton P., Jimenez S.A., Prockop D.J.;  
 RT "Spondyloepiphyseal dysplasia and precocious osteoarthritis in a  
 RT family with an Arg75->Cys mutation in the procollagen type II gene  
 RT (COL2A1).";  
 RL Hum. Genet. 92:499-505(1993).  
 RN [24]  
 RP VARIANT SEDC CYS-920.  
 RX MEDLINE-93315508; PubMed-8325695;  
 RA Chan D., Taylor T.K.F., Cole W.G.;  
 RT "Characterization of an arginine 789 to cysteine substitution in  
 RT alpha 1 (II) collagen chains of a patient with spondyloepiphyseal  
 RT dysplasia.";  
 RL J. Biol. Chem. 268:15238-15245(1993).  
 RN [25]  
 RP VARIANT SEDC SER-1128.  
 RX MEDLINE-93140139; PubMed-8423604;  
 RA Cole W.G., Hall R.K., Rogers J.G.;  
 RT "The clinical features of spondyloepiphyseal dysplasia congenita  
 RT resulting from the substitution of glycine 997 by serine in the alpha  
 RT 1(II) chain of type II collagen.";  
 RL J. Med. Genet. 30:27-35(1993).

Query Match 88.8%; Score 71; DB 1; Length 1418;  
 Best Local Similarity 80.0%; Pred. No. 0.0049;  
 Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0.

QY 1 GTPGPGGIAGGGRVY 15  
 Db 900 GPGPGGIAGGGRIV 914  
 I|||||:||||:|

RESULT 7  
 CA12\_MOUSE  
 ID CA12\_MOUSE STANDARD: PRT: 1459 AA.  
 AC P28481;  
 DT 01-DEC-1992 (Rel. 24, Created)  
 DT 01-DEC-1992 (Rel. 24, Last sequence update)  
 DT 28-FEB-2003 (Rel. 41, Last annotation update)  
 DE Collagen alpha 1(II) chain precursor [contains: Chondrocalcin].  
 GN COL2A1.  
 OS Mus musculus (Mouse).  
 CC Eukaryota; Euteleostomi; Chordata; Vertebrata; Euteleostomi;  
 CC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
 OX NCBI\_TaxID=10090;  
 RX MEDLINE-91358489; PubMed-1885613;  
 RA Metastanta M., Toman D., de Crombrughe B., Vuorio E.;

RT	"Mouse type II collagen gene. Complete nucleotide sequence, exon
RT	structure, and alternative splicing."
RL	J. Biol. Chem. 266:16862-16869(1991).
RN	[2]
RP	SEQUENCE OF 1455-1459 FROM N.A.
RA	MEDLINE=91274355; PubMed=205484;
RX	Metsaranta M., Toman D., de Crombrughe B., Vuorio E.:
RT	"Specific hybridization probes for mouse type I, II, III and IX
RT	collagen mRNAs."
RL	Biochim. Biophys. Acta 1089:241-243(1991).
CC	-I- FUNCTION: COLLAGEN TYPE II IS SPECIFIC FOR CARTILAGINOUS TISSUES.
CC	-I- SUBUNIT: TRIMERS OF IDENTICAL ALPHA 1(II) CHAINS.
CC	-I- ALTERNATIVE PRODUCTS:
CC	Event-Alternative splicing; Named isoforms=2;
CC	Name-Long:
CC	Isoid=P28481-1; Sequence=Displayed;
CC	Name-Short:
CC	Isoid=P28481-2; Sequence=VSP_001139, VSP_001140;
CC	-I- PPM: Prolines at the third position of the tripeptide repeating
CC	unit (G-X-Y) are hydroxylated in some or all of the chains.
CC	-I- SIMILARITY: Contains 1 WMC domain.
CC	-----
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CC	or send an email to <a href="mailto:license@isb-sib.ch">license@isb-sib.ch</a> ).
CC	-----
DR	EMBL; M65161; AAA68100.1; -.
DR	EMBL; X57982; CAA41047.1; -.
DR	MGD; MG1; 88452; Col2a1.
DR	InterPro: IPRO00087; Collagen.
DR	InterPro: IPRO00085; Fib.collagen_C.
DR	InterPro: IPRO01007; WFC.
DR	Pfam; PF01410; COLF1; 1.
DR	Pfam; PF01391; Collagen; 18.
DR	Pfam; PF00093; WVC; 1.
DR	ProdDom: PD000007; Clq_hellix; 5.
DR	ProdDom: PD002078; Fib.collagen_C; 1.
DR	SMART; SM00214; WVC; 1.
DR	SMART; SM01208; WFC_1; 1.
DR	PROSITE; PS01084; WFC_2; 1.
KW	Extracellular matrix; Connective tissue; Repeat; Hydroxylation;
KW	Glycoprotein; Collagen; Cartilage; Signal; Alternative splicing.
FT	SIGNAL 1 25
FT	PROPEP 26 153
FT	AMINO-TERMINAL PROPEPTIDE
FT	(BY SIMILARITY).
FT	COLLAGEN ALPHA 1(II) CHAIN.
FT	CHONDROCALCIN.
FT	WFC.
FT	TRIPLE-HELICAL REGION.
FT	NONHELIICAL REGION (C-TERMINAL).
FT	O -> R (in isoform Short).
FT	/FTid-VSP_001139.
FT	VARSPLIC 29 29
FT	Missing (in isoform Short).
FT	/FTid-VSP_001140.
FT	VARSPPLIC 30 98
FT	Missing (in isoform Short).
SO	SEQUENCE 1459 AA; 139154 MW; F6C84FA7C532E/F2 CRC64;
QY	Query Match 88.8%; Score 71; DB 1; Length 1459;
Db	Best Local Similarity 80.0%; Pred. No. 0.0051;
Matches	12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
ID	CA24_ASCSU STANDARD; PRT; 1763 AA.
RESULT 8	
CA24_ASCSU	
ID	CA24_ASCSU

AC P27393; 01-AUG-1992 (Rel. 23, Created)  
 DT 01-AUG-1992 (Rel. 23, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Collagen alpha 2(IV) chain precursor.  
 OS Ascaris suum (Pig roundworm) (Ascaris lumbricoides).  
 OC Eukaryota; Metazoa; Nematoda; Chromadorea; Ascaridida; Ascaridoidea;  
 OC Ascaridae; Ascaris.  
 NCBI\_TaxID=6253;  
 RX MEDLINE=91340768; PubMed=1714907;  
 RA Pettitt J., Kingston I.B.,  
 RT "The complete primary structure of a nematode alpha 2(IV) collagen  
 and the partial structural organization of its gene."  
 RL J. Biol. Chem. 266:16149-16156(1991).  
 CC -I- FUNCTION: COLLAGEN TYPE IV IS SPECIFIC FOR BASEMENT MEMBRANS.  
 CC -I- SUBUNIT: TRIMERS OF TWO ALPHA 1(IV) AND ONE ALPHA 2(IV) CHAIN.  
 CC TYPE IV COLLAGEN FORMS A MESH-LIKE NETWORK LINKED THROUGH  
 CC INTERMOLECULAR INTERACTIONS BETWEEN 7S DOMAINS AND BETWEEN NCI  
 CC DOMAINS.  
 CC -I- ALTERNATIVE PRODUCTS:  
 CC Event-Alternative splicing. Named isoforms=2;  
 CC Name=I;  
 CC IsoId=P27393-1; Sequence=Displayed;  
 CC Name=II;  
 CC IsoId=P27393-2; Sequence=VSP\_001159;  
 CC -I- DOMAIN: ALPHA CHAINS OF TYPE IV COLLAGEN HAVE A NONCOLLAGENOUS  
 CC DOMAIN (NC1) AT THEIR C-TERMINUS. FREQUENT INTERRUPTIONS OF THE  
 CC G-X-Y REPEATS IN THE LONG CENTRAL TRIPLE-HELICAL DOMAIN (WHICH MAY  
 CC CAUSE FLEXIBILITY IN THE TRIPLE HELIX), AND A SHORT N-TERMINAL  
 CC TRIPLE-HELICAL 7S DOMAIN.  
 CC -I- PWM: Prolines at the third position of the tripeptide repeating  
 CC unit (G-X-Y) are hydroxylated in some or all of the chains.  
 CC -I- PWM: Type IV collagens contain numerous cysteine residues which  
 CC are involved in inter- and intramolecular disulfide bonding. 12 of  
 CC these, located in the NC1 domain, are conserved in all known type  
 CC IV collagens.  
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 CC or send an email to [license@sib-sib.ch](mailto:license@sib-sib.ch)).  
 CC -----  
 CC DR EMBL; M67507; AAA18014.1; -;  
 CC PIR; S16366;  
 CC InterPro; IPR000087; Collagen.  
 CC InterPro; IPR001442; ProcollagenC4.  
 CC Pfam; PF01413; C4; 2.  
 CC Pfam; PF01391; Collagen; 25.  
 CC ProDom; PD000007; Ctg helix; 6.  
 CC ProDom; PD003923; ProcollagenC4; 1.  
 CC SMART; SM00111; C4; 2.  
 CC DR Hydroxylation; Connective tissue; Basement membrane; Repeat; Collagen;  
 CC Alternative splicing; Glycoprotein; Signal.  
 CC FT SIGNAL 1 26  
 CC FT CHAIN 27 1763 COLLAGEN ALPHA 2(IV) CHAIN.  
 CC FT DOMAIN 27 42  
 CC FT DOMAIN 43 1529 TRIPLE-HELICAL REGION.  
 CC FT DOMAIN 1530 1763 NONHELICAL REGION (NC1).  
 CC FT DISULFID 1548 1637 OR 1634 (BY SIMILARITY).  
 CC FT DISULFID 1581 1634 OR 1637 (BY SIMILARITY).  
 CC FT DISULFID 1593 1599 BY SIMILARITY.  
 CC FT DISULFID 1656 1752 OR 1749 (BY SIMILARITY).  
 CC FT DISULFID 1690 1749 OR 1752 (BY SIMILARITY).  
 CC FT DISULFID 1702 1709 BY SIMILARITY.  
 CC FT CARBOHYD 126 126 N-LINKED (GLCNAC. . .) (POTENTIAL).  
 CC FT CARBOHYD 249 249 O-LINKED (XYL. . .) (GLYCOSAMINOGLYCAN)  
 CC FT (IN ISOFORM II) (POTENTIAL).  
 CC FT VARSPLIC 230 266 GEOGPRGQGPGRPVSTGAGCTITGEGAGMGK ->

FT FT GDICGAPGPPGPPRFTSGSIVGRGHSQDKGVK (in  
 FT FT Isoform II).  
 FT FT /FTId=VSP\_001159.  
 SQ SEQUENCE 1763 AA; 168526 MW; 304F52B8C06AA80D CRC64;  
 Query Match 77.5%; Score 62; DB 1; Length 1763;  
 Best Local Similarity 66.7%; Pred. No. 0.14;  
 Matches 10; Conservative 2; Mismatches 3; Indels 0; Gaps 0;  
 QY 1 GTPGPGIAGGQGV 15  
 Db 82 GPPGPGIKGRGRT 96  
 RESULT 9  
 CA29\_HUMAN STANDARD; PRT; 689 AA.  
 AC Q14055;  
 DT 28-FEB-2003 (Rel. 41, Created)  
 DT 28-FEB-2003 (Rel. 41, Last sequence update)  
 DE 28-FEB-2003 (Rel. 41, Last annotation update)  
 DE Collagen alpha 2(IX) chain precursor.  
 GN COL9A2.  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 OC NCBI\_TaxID=9606;  
 CC [1]  
 CC SEQUENCE FROM N.A.  
 CC TISSUE-Foreskin;  
 CC MEDLINE=98370844; PubMed=9707347;  
 CC RA Pihlajamaa T., Vuorio M.M., Annunen S., Peraelae M., Prockop D.J.,  
 CC Ala-Kokko L.;  
 CC "Human COL9A1 and COL9A2 genes. Two genes of 90 and 15 kb code for  
 CC similar polypeptides of the same collagen molecule."  
 CC RT Matrix Biol. 17:237-241(1998).  
 CC [2]  
 CC SEQUENCE FROM N.A.  
 CC RA Donnelly S.;  
 CC RL submitted (FEB-2000) to the EMBL/GenBank/DBJ databases.  
 CC [3]  
 CC SEQUENCE OF 50-668 FROM N.A.  
 CC RC TISSUE-Cartilage;  
 CC RX MEDLINE=9320262; PubMed=8454052;  
 CC RA Peraelae M., Hanninen M., Hasbicka J., Elima K., Vuorio E.;  
 CC RT "Molecular cloning of the human alpha 2(IX) collagen cDNA and  
 CC assignment of the human COL9A2 gene to chromosome 1.";  
 CC RL FEBS Lett. 319:177-180(1993).  
 CC [4]  
 CC VARIANT IDD TRP-326, AND VARIANT ARG-326.  
 CC RX MEDLINE=99340300; PubMed=10411504;  
 CC RA Annunen S., Paasilta P., Lohiniva J., Peraelae M., Pihlajamaa T.,  
 CC RA Karppinen J., Tervonen O., Kroeger H., Laehte S., Vanharanta H.,  
 CC RA Ryhanen L., Goering H.H., Ott J., Prockop D.J., Ala-Kokko L.;  
 CC "An allele of COL9A2 associated with intervertebral disc disease."  
 CC RT Science 285:409-412(1999).  
 CC CC -I- FUNCTION: Structural component of hyaline cartilage and vitreous  
 CC of the eye.  
 CC CC -I- SUBUNIT: Heterotrimer of an alpha 1(IX), an alpha 2(IX) and an  
 CC alpha 3(IX) chain.  
 CC CC -I- PWM: Covalently linked to the telopeptides of type II collagen by  
 CC lysine-derived cross-links.  
 CC CC -I- PWM: Prolines at the third position of the tripeptide repeating  
 CC unit (G-X-Y) are hydroxylated in some or all of the chains.  
 CC CC -I- DISEASE: Defects in COL9A2 are a cause of multiple epiphyseal  
 CC dysplasia type 2 (MED2) [MIM:600204]. MED2 is characterized by  
 CC flattened, irregular epiphyses in most joints, particularly the  
 CC knees. In childhood, typical features include waddling gait and  
 CC knee pain/stiffness. Inheritance is autosomal dominant.  
 CC CC -I- DISEASE: Defects in COL9A2 are a cause of susceptibility to  
 CC intervertebral disc disease (IDD) [MIM:603932], one of the  
 CC most common musculo-skeletal disorders.  
 CC CC -I- SIMILARITY: BELONGS TO THE FIBRIL-ASSOCIATED COLLAGENS WITH

----- INTERRUPTED HELICES (FACIT) FAMILY. -----

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CC -----

DR EMBL: AF019406; AAC33512.1; -  
 DR EMBL: AL050341; CAB81611.1; -  
 DR EMBL: M95610; AAA80977.1; -  
 DR Genew: HGNC:2218; COL9A2.  
 DR MIM: 120260; -  
 DR MIM: 600204; -  
 DR MIM: 603932; -  
 DR GO: GO:0005594; C:collagen type IX; TAS.  
 DR GO: GO:0005202; F:collagen; TAS.  
 DR GO: GO:0001501; F:skeletal development; TAS.  
 DR InterPro: IPR000087; Collagen.  
 DR Pfam: PF01391; Collagen; 9.  
 DR Prodom: PD000007; Ctg helix; 1.  
 KW Extracellular matrix; Connective tissue; Repeat; Hydroxylation;  
 KW Cartilage; Collagen; Signal; Disease mutation; Polymorphism.  
 KW SIGNAL 1 23  
 FT CHAIN 1 23  
 FT DOMAIN 24 689 COLLAGEN ALPHA 2(IX) CHAIN.  
 FT DOMAIN 27 519 TRIPLE-HELICAL REGION 3 (COL3).  
 FT DOMAIN 520 549 NONHELICAL REGION 3 (NC3).  
 FT DOMAIN 550 632 TRIPLE-HELICAL REGION 2 (COL2).  
 FT DOMAIN 633 634 NONHELICAL REGION 2 (NC2).  
 FT DOMAIN 635 664 TRIPLE-HELICAL REGION 1 (COL1).  
 FT DOMAIN 665 689 NONHELICAL REGION 1 (NC1).  
 FT VARIANT 326 326 Q -> R.  
 FT VARIANT 326 326 /FTID=VAR.012659.  
 FT VARIANT 326 326 Q -> W (in IDD; requires 2 nucleotide  
 FT substitutions).  
 FT /FTID=VAR.012658.  
 SQ SEQUENCE 689 AA: 65131 MW: EB6106E02F6FA862 CRC64:

Query Match 73.8%; Score 59; DB 1; Length 689;  
 Best Local Similarity 71.4%; Pred. No. 0.16;  
 Matches 10; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGV 14  
 | | | | | | | | | |  
 Db 493 GPGPGRIAGNRGV 506

----- RESULT 10 -----

CA21\_CHICK STANDARD: PRT: 1362 AA.  
 AC P02457; P87491; P87492; Q90758; Q90792; Q90795; Q90797; Q92014;  
 DT 21-JUL-1986 (Rel. 01, Created)  
 DT 15-DEC-1998 (Rel. 37, Last sequence update)  
 DT 28-FEB-2003 (Rel. 41, Last annotation update)  
 DE Collagen alpha 2(I) chain precursor (Fragments).  
 GN COL1A2.  
 OS Gallus gallus (Chicken).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Archosauria; Aves; Neognathae; Galliformes; Phasianidae;  
 OC Gallus.  
 OC NCBI\_TaxID=9031;  
 RN [1]  
 RP SEQUENCE OF 1-245: 262-448 AND 466-1362 FROM N.A.  
 RX MEDLINE=86185168; PubMed=3868961;  
 RA Boedtker H., Finer M., Aho S.;  
 RL Ann. N.Y. Acad. Sci. 460:85-116(1985).  
 RN [2]  
 RP SEQUENCE OF 1-89 FROM N.A.  
 RX MEDLINE=83246518; PubMed=6135195;  
 RA Tate V.E., Finer M.H., Boedtker H., Doty P.;

RT "Chick pro alpha 2 (I) collagen gene: exon location and coding  
 RT potential for the prepropeptide."  
 RL Nucleic Acids Res. 11:91-104(1983).  
 RN [3]  
 RP SEQUENCE OF 1-14 FROM N.A.  
 RX MEDLINE=82060240; PubMed=6946474;  
 RA Vogel I.G., Ohkubo H., Sobel M.E., Yamada Y., Pastan I.,  
 RA de Crombrughe B.;  
 RT "Structure of the promoter for chicken alpha 2 type I collagen gene."  
 RL Proc. Natl. Acad. Sci. U.S.A. 78:5334-5338(1981).  
 RN [4]  
 RP SEQUENCE OF 1-33 FROM N.A.  
 RX MEDLINE=84297217; PubMed=6473103;  
 RA Aho S., Tate V.E., Boedtker H.;  
 RT "Location of the 11 bp exon in the chicken pro alpha 2(I) collagen  
 RT gene."  
 RL Nucleic Acids Res. 12:6117-6125(1984).  
 RN [5]  
 RP SEQUENCE OF 1-79 FROM N.A.  
 RX MEDLINE=88056316; PubMed=3678834;  
 RA Finer M.H., Boedtker H., Doty P.;  
 RT "Construction and characterization of cDNA clones encoding the 5' end  
 RT of the chicken pro alpha 1(I) collagen mRNA."  
 RL Gene 56:71-78(1987).  
 RN [6]  
 RP SEQUENCE OF 78-92.  
 RC TISSUE=Skin;  
 RX MEDLINE=7115216; PubMed=5544653;  
 RA Hightberger J.H., Kang A.H., Gross J.;  
 RT "Comparative studies on the amino acid sequence of the alpha 2-CB2  
 RT peptides from chick and rat skin collagens."  
 RL Biochemistry 10:610-616(1971).  
 RN [7]  
 RP SEQUENCE OF 74-91; 263-448 AND 1088-1169 FROM N.A.  
 RX MEDLINE=82058081; PubMed=6272119;  
 RA Wozney J., Hanahan D., Tate V.E., Boedtker H., Doty P.;  
 RT "Structure of the pro alpha 2 (I) collagen gene."  
 RL Nature 294:129-135(1981).  
 RN [8]  
 RP SEQUENCE OF 78-92.  
 RC TISSUE=Skin;  
 RX MEDLINE=70131186; PubMed=4313735;  
 RA Kang A.H., Gross J.;  
 RT "Amino acid sequence of cyanogen bromide peptides from the amino-  
 RT terminal region of chick skin collagen."  
 RL Biochemistry 9:796-804(1970).  
 RN [9]  
 RP SEQUENCE OF 78-92 AND 415-448.  
 RC TISSUE=Skin;  
 RX MEDLINE=69285369; PubMed=5809220;  
 RA Kang A.H., Igarashi S., Gross J.;  
 RT "Characterization of the cyanogen bromide peptides from the alpha-2  
 RT chain of chick skin collagen."  
 RL Biochemistry 8:3200-3204(1969).  
 RN [10]  
 RP SEQUENCE OF 78-92 AND 415-448.  
 RC TISSUE=Bone;  
 RX MEDLINE=69206882; PubMed=5785233;  
 RA Lane J.M., Miller E.J.;  
 RT "Isolation and characterization of the peptides derived from the  
 RT alpha 2 chain of chick bone collagen after cyanogen bromide  
 RT cleavage."  
 RL Biochemistry 8:2134-2139(1969).  
 RN [11]  
 RP SEQUENCE OF 566-587 FROM N.A.  
 RX MEDLINE=79074829; PubMed=364479;  
 RA Lehman H., Frischau A.-M., Hanahan D., Wozney J., Fuller F.,  
 RA Trehanjankov R., Boedtker H., Doty P.;  
 RT "Construction and characterization of a 2.5-Kilobase procollagen  
 RT clone."  
 RL Proc. Natl. Acad. Sci. U.S.A. 75:5417-5421(1978).  
 RN [12]  
 RP SEQUENCE OF 902-1362 FROM N.A.

RX MEDLINE=81160715; PubMed=6927845;  
 RA Fuller F., Boedtker H.;  
 RT "Sequence determination and analysis of the 3' region of chicken pro-  
 alpha 1(I) and pro-alpha 2(I) collagen messenger ribonucleic acids  
 including the carboxy-terminal propeptide sequences.";  
 RL Biochemistry 20:996-1006(1981).  
 RN [13]  
 RP SEQUENCE OF 998-1169 AND 1234-1362 FROM N.A.  
 RX MEDLINE=81264246; PubMed=6267043;  
 RA Dickson L.A., Ninomiya Y., Bernard M.P., Pesciotta D.M., Parsons J.,  
 RA Green G., Eikenberry E.F., de Crombrughe B., Vogeli G., Pastan I.,  
 RA Flatzak P.P., Olsen B.R.;  
 RT "The exon/intron structure of the 3'-region of the pro alpha 2(I)  
 collagen gene.";  
 RL J. Biol. Chem. 256:8407-8415(1981).  
 RN [14]  
 RP SEQUENCE OF 932-954 AND 968-980 FROM N.A.  
 RX MEDLINE=81064671; PubMed=6159982;  
 RA Avvedimento V.E., Vogeli G., Yamada Y., Maizel J.V. Jr., Pastan I.,  
 RA de Crombrughe B.;  
 RT "Correlation between splicing sites within an intron and their  
 sequence complementarity with U1 RNA.";  
 RL Cell 21:689-696(1980).  
 RN [15]  
 RP SEQUENCE OF 126-161, 467-517 AND 926-954 FROM N.A.  
 RX MEDLINE=8112157; PubMed=7460017;  
 RA Yamada Y., Avvedimento V.E., Mudry M., Ohkubo H., Vogeli G.,  
 RA Imani M., Pastan I., de Crombrughe B.;  
 RT "The collagen gene: evidence for its evolutionary assembly by  
 amplification of a DNA segment containing an exon of 54 bp.";  
 RL Cell 22:887-892(1980).  
 CC -1- FUNCTION: TYPE I COLLAGEN IS A MEMBER OF GROUP I COLLAGEN  
 CC (FIBRILLAR FORMING COLLAGEN).  
 CC -1- SUBUNIT: TRIMERS OF ONE ALPHA 2(I) AND TWO ALPHA 1(I) CHAINS.  
 CC -1- TISSUE SPECIFICITY: FORMS THE FIBRILS OF TENDON, LIGAMENTS AND  
 CC BONES. IN BONES THE FIBRILS ARE MINERALIZED WITH CALCIUM  
 CC HYDROXYAPATITE.  
 CC -1- PTM: Prolines at the third position of the tripeptide repeating  
 CC unit (G-X-Y) are hydroxylated in some or all of the chains.  
 CC -----  
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 CC -----  
 DR EMBL; M25963; AAA69960.1; -;  
 DR EMBL; M25956; AAA69960.1; JOINED.  
 DR EMBL; M25959; AAA69960.1; JOINED.  
 DR EMBL; M25961; AAA69960.1; JOINED.  
 DR EMBL; M25962; AAA69960.1; JOINED.  
 DR EMBL; M25965; AAA69961.1; -;  
 DR EMBL; M25964; AAA69961.1; JOINED.  
 DR EMBL; M25984; AAA69962.1; -;  
 DR EMBL; M25957; AAA69962.1; JOINED.  
 DR EMBL; M25966; AAA69962.1; JOINED.  
 DR EMBL; M25967; AAA69962.1; JOINED.  
 DR EMBL; M25969; AAA69962.1; JOINED.  
 DR EMBL; M25970; AAA69962.1; JOINED.  
 DR EMBL; M25971; AAA69962.1; JOINED.  
 DR EMBL; M25972; AAA69962.1; JOINED.  
 DR EMBL; M25973; AAA69962.1; JOINED.  
 DR EMBL; M25974; AAA69962.1; JOINED.  
 DR EMBL; M25976; AAA69962.1; JOINED.  
 DR EMBL; M25977; AAA69962.1; JOINED.  
 DR EMBL; M25978; AAA69962.1; JOINED.  
 DR EMBL; M25979; AAA69962.1; JOINED.  
 DR EMBL; M25980; AAA69962.1; JOINED.  
 DR EMBL; M25981; AAA69962.1; JOINED.  
 DR EMBL; M25982; AAA69962.1; JOINED.  
 DR EMBL; M25983; AAA69962.1; JOINED.

DR EMBL; J00826; AAA51611.1; -;  
 DR EMBL; J00821; AAA51611.1; JOINED.  
 DR EMBL; K00792; AAA51611.1; JOINED.  
 DR EMBL; J00830; AAA51613.1; -;  
 DR EMBL; J00829; AAA51613.1; JOINED.  
 DR EMBL; J00837; AAA51614.1; -;  
 DR EMBL; J00812; AAA51615.1; -;  
 DR EMBL; J00811; AAA51615.1; JOINED.  
 DR EMBL; J00814; AAA51615.1; JOINED.  
 DR EMBL; J00815; AAA51615.1; JOINED.  
 DR EMBL; X02557; CAA26493.1; -;  
 DR EMBL; K00794; -; NOT\_ANNOTATED\_CDS.  
 DR EMBL; V00390; CAA23688.1; -;  
 DR EMBL; M17608; AAA48673.1; -;  
 DR EMBL; M10581; AAA48637.1; -;  
 DR EMBL; M10540; AAA48638.1; -;  
 DR EMBL; J00828; AAA51612.1; -;  
 DR EMBL; J00827; AAA51612.1; JOINED.  
 DR EMBL; J00831; -; NOT\_ANNOTATED\_CDS.  
 DR EMBL; J00832; -; NOT\_ANNOTATED\_CDS.  
 DR EMBL; J00833; -; NOT\_ANNOTATED\_CDS.  
 DR EMBL; J00822; -; NOT\_ANNOTATED\_CDS.  
 DR PIR; I50173; I50173.  
 DR PIR; I50206; CGCH2S.  
 DR InterPro; IPR000087; Collagen.  
 DR InterPro; IPR000885; Fib\_Collagen\_C.  
 DR Pfam; PF01391; Collagen; 17.  
 DR Pfam; PF01410; COLTR; 1.  
 DR ProDom; PD000007; Ctg\_helix; 4.  
 DR ProDom; PD002078; Fib\_collagen\_C; 1.  
 DR SMART; SM00038; COLFI; 1.  
 KW Extracellular matrix; Connective tissue; Repeat; Hydroxylation;

Query Match 72.5%; Score 58; DB 1; Length 1362;  
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 Qy 1 GTPGPGIAGGQGVV 15  
 Db 857 GTPGPGIAGGQGVV 871

RESULT 11  
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 AC P05997;  
 DT 01-APR-1988 (Rel. 07, Created)  
 DT 01-JAN-1990 (Rel. 13, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Collagen alpha 2(V) chain precursor.  
 GN COL5A2.  
 OS Homo sapiens (Human).  
 CC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 CC Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.  
 OX NCBI\_Taxid=9606;  
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 RP SEQUENCE OF 1-463 FROM N.A.  
 RX MEDLINE=89123368; PubMed=2914927;  
 RA Woodbury D., Benson-Chanda V., Ramirez F.;  
 RT "Amino-terminal propeptide of human pro-alpha 2(V) collagen conforms  
 RT to the structural criteria of a fibrillar procollagen molecule.";  
 RL J. Biol. Chem. 264:2735-2738(1989).  
 RN [2]  
 RP SEQUENCE OF 398-1496 FROM N.A.  
 RX MEDLINE=87146331; PubMed=3029669;  
 RA Weil D., Bernard M.P., Gargano S., Ramirez F.;  
 RT "The pro alpha 2(V) collagen gene is evolutionarily related to the  
 RT major fibrillar-forming collagens.";  
 RL Nucleic Acids Res. 15:181-198(1987).  
 RN [3]  
 RP SEQUENCE OF 1227-1496 FROM N.A.  
 RX MEDLINE=85289337; PubMed=2411731;  
 RA Myers J.C., Loidl H.R., Seyer J.M., Dion A.S.;



RN [2]  
 RP SEQUENCE OF 73-636 FROM N.A.  
 RC STRAIN-Sprague-Dawley; TISSUE=Fibroblast;  
 RA Muttz T., Ellerstrom C., Lundmark C., Christenson C.;  
 RL Submitted (APR-1998) to the EMBL/Genbank/DBJ databases.  
 RN [3]  
 RP SEQUENCE OF 308-482 FROM N.A.  
 RX MEDLINE=88296083; PubMed=2456904;  
 RA Frankel F.R., Hsu C.-Y.J., Meyers J.C., Lin E., Lytle C.R.,  
 RA Komm B., Mohr K.;  
 RT "Regulation of alpha 2(I), alpha 1(III), and alpha 2(V) collagen  
 RT mRNAs by estradiol in the immature rat uterus.";  
 RL DNA 7,347-354(1988).  
 CC -1- FUNCTION: COLLAGEN TYPE III OCCURS IN MOST SORT CONNECTIVE TISSUES  
 CC ALONG WITH TYPE I COLLAGEN.  
 CC -1- SUBUNIT: TRIMERS OF IDENTICAL ALPHA 1(III) CHAINS. THE CHAINS ARE  
 CC LINKED TO EACH OTHER BY INTERCHAIN DISULFIDE BONDS. TRIMERS ARE  
 CC ALSO CROSS-LINKED VIA HYDROXYLINES.  
 CC -1- PTM: Prolines at the third position of the tripeptide repeating  
 CC unit (G-X-Y) are hydroxylated in some or all of the chains.  
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 CC -----  
 DR EMBL: X70369; CAA49832.1; -;  
 DR EMBL: AJ005395; CAA06510.1; -;  
 DR EMBL: M21354; AAA40942.1; -;  
 DR PIR: S41067; S41067.  
 DR InterPro: IPR000087; Collagen.  
 DR InterPro: IPR000885; Fib\_collagen\_C.  
 DR InterPro: IPR001007; VWF\_C.  
 DR Pfam: PF01410; COLF1; 1.  
 DR Pfam: PF01391; Collagen; 6.  
 DR ProDom: PD000007; C1g\_helix; 1.  
 DR ProDom: PD002078; Fib\_collagen\_C; 1.  
 DR SMART: SMO0038; COLF1; 1.  
 DR PROSITE: PS01208; VWF\_C\_1; PARTIAL.  
 DR Extracellular matrix; Connective tissue; Repeat; Hydroxylation;  
 KW Collagen; Glycoprotein.  
 FT CHAIN  
 FT NON\_TER 1 1  
 FT PROPEP 376 636 COLLAGEN ALPHA 1(III) CHAIN.  
 FT DOMAIN 369 636 CARBOXYL-TERMINAL PROPEPTIDE.  
 FT DISULFID 368 636 TRIPLE-HELICAL REGION.  
 FT DISULFID 369 636 NONHELICAL REGION (C-TERMINAL).  
 FT CONFLICT 340 340 INTERCHAIN (BY SIMILARITY).  
 FT CONFLICT 429 429 N -> D (IN REF. 2).  
 FT CONFLICT 429 429 A -> G (IN REF. 2).  
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 Best Local Similarity 69.2%; Pred. No. 0.3;  
 Matches 9; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OS Bos taurus (Bovine).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;  
 OC Bovidae; Bovinae; Bos.  
 OC NCBI\_TaxID=9913;  
 RN [1]  
 RP SEQUENCE OF 1-242.  
 RX MEDLINE=80026026; PubMed=488906;  
 RA Fietzek P.P., Allmann H., Rautenberg J., Henkel W., Wächter E.,  
 RA Kuhn K.;  
 RT "The covalent structure of calf skin type III collagen. I. The amino  
 RT acid sequence of the amino terminal region of the alpha 1(III) chain  
 RT (positions 1-222).";  
 RL Hoppe-Seyler's Z. Physiol. Chem. 360:809-820(1979).  
 RN [2]  
 RP SEQUENCE OF 243-422.  
 RX MEDLINE=80026027; PubMed=488907;  
 RA Dewes H., Fietzek P.P., Kuhn K.;  
 RT "The covalent structure of calf skin type III collagen. II. The amino  
 RT acid sequence of the cyanogen bromide peptide alpha 1(III)CB1,8,10,2  
 RT (positions 223-402).";  
 RL Hoppe-Seyler's Z. Physiol. Chem. 360:821-832(1979).  
 RN [3]  
 RP SEQUENCE OF 423-571.  
 RX MEDLINE=80026028; PubMed=488908;  
 RA Bentz H., Fietzek P.P., Kuhn K.;  
 RT "The covalent structure of calf skin type III collagen. III. The  
 RT amino acid sequence of the cyanogen bromide peptide alpha 1(III)CB4  
 RT (positions 403-551).";  
 RL Hoppe-Seyler's Z. Physiol. Chem. 360:833-840(1979).  
 RN [4]  
 RP SEQUENCE OF 572-808.  
 RX MEDLINE=80026029; PubMed=488909;  
 RA Lang H., Glanville R.W., Fietzek P.P., Kuhn K.;  
 RT "The covalent structure of calf skin type III collagen. IV. The amino  
 RT acid sequence of the cyanogen bromide peptide alpha 1(III)CB5  
 RT (positions 552-788).";  
 RL Hoppe-Seyler's Z. Physiol. Chem. 360:841-850(1979).  
 RN [5]  
 RP SEQUENCE OF 809-947.  
 RX MEDLINE=80026030; PubMed=488910;  
 RA Dewes H., Fietzek P.P., Kuhn K.;  
 RT "The covalent structure of calf skin type III collagen. V. The amino  
 RT acid sequence of the cyanogen bromide peptide alpha 1(III)CB9A  
 RT (position 789-927).";  
 RL Hoppe-Seyler's Z. Physiol. Chem. 360:851-860(1979).  
 RN [6]  
 RP SEQUENCE OF 948-1049.  
 RX MEDLINE=80026031; PubMed=488911;  
 RA Allmann H., Fietzek P.P., Glanville R.W., Kuhn K.;  
 RT "The covalent structure of calf skin type III collagen. VI. The amino  
 RT acid sequence of the carboxyterminal cyanogen bromide peptide alpha  
 RT 1(III)CB9B (positions 928-1028).";  
 RL Hoppe-Seyler's Z. Physiol. Chem. 360:861-868(1979).  
 CC -1- FUNCTION: COLLAGEN TYPE III OCCURS IN MOST SORT CONNECTIVE TISSUES  
 CC ALONG WITH TYPE I COLLAGEN.  
 CC -1- SUBUNIT: TRIMERS OF IDENTICAL ALPHA 1(III) CHAINS. THE CHAINS ARE  
 CC LINKED TO EACH OTHER BY INTERCHAIN DISULFIDE BONDS. TRIMERS ARE  
 CC ALSO CROSS-LINKED VIA HYDROXYLINES.  
 CC -1- PTM: Prolines at the third position of the tripeptide repeating  
 CC unit (G-X-Y) are hydroxylated in some or all of the chains.  
 CC -----  
 DR PIR: A02862; CGB075.  
 DR InterPro: IPR000087; Collagen.  
 DR InterPro: IPR001007; VWF\_C.  
 DR Pfam: PF01391; Collagen; 17.  
 DR ProDom: PD000007; C1g\_helix; 3.  
 DR PROSITE: PS01208; VWF\_C\_1; PARTIAL.  
 DR Extracellular matrix; Connective tissue; Repeat; Hydroxylation;  
 KW Glycoprotein; Collagen.  
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 FT NON\_TER 1 14  
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 FT DOMAIN 1041 1049 TRIPLE-HELICAL REGION.  
 FT MOD\_RES 95 95 NONHELICAL REGION (C-TERMINAL).  
 FT HYDROXYLATION.

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FT MOD_RES 107 107 HYDROXYLATION.
FT MOD_RES 119 119 HYDROXYLATION.
FT MOD_RES 938 938 HYDROXYLATION.
FT MOD_RES 950 950 HYDROXYLATION.
FT CARBOHYD 107 107 O-LINKED (GAL. . .).
FT CARBOHYD 950 950 O-LINKED (GAL. . .).
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FT DISULFID 1041 1041 INTERCHAIN.
SQ SEQUENCE 1049 AA; 93651 MW; 8BEC33D1C66EC9A3 CRC64;

Query Match 71.2%; Score 57; DB 1; Length 1049;
Best Local Similarity 69.2%; Pred. No. 0.47;
Matches 9; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 GTPGPGIAGGRC 13
Db 696 GPPGPGVKGRC 708

RESULT 14
CA13_CHICK STANDARD; PRT: 1262 AA.
AC P12105; P79758; P79759; Q90794; Q92029;
DT 01-OCT-1989 (Rel. 12, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 28-FEB-2003 (Rel. 41, Last annotation update)
DE Collagen alpha 1(III) chain precursor (Fragments).
GN COL3A1.
OS Gallus gallus (Chicken).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Atherosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
OC Gallus.
OX NCBI_TaxID=9031;
RN [1]
RP SEQUENCE OF 1-886 FROM N.A.
RC TISSUE=Kidney;
RX MEDLINE=94266842; PubMed=8206952;
RA Nah H.-D., Niu Z., Adams S.L.;
RT "An alternative transcript of the chick type III collagen gene that
RT does not encode type III collagen.";
RN [2]
RP SEQUENCE OF 29-96; 332-397; 431-484; 503-535 AND 869-976 FROM N.A.
RX MEDLINE=84270696; PubMed=6547770;
RA Yamada Y., Lieu G., Mudryj M., Obici S., de Crombrughe B.;
RT "Conservation of the sites for one but not another class of exons in
RT two chick collagen genes.";
RN [3]
RP SEQUENCE OF 977-1262 FROM N.A.
RX MEDLINE=83220816; PubMed=6856474;
RA Yamada Y., Kuhn K., de Crombrughe B.;
RT "A conserved nucleotide sequence, coding for a segment of the C-
RT propeptide, is found at the same location in different collagen
RT genes.";
RN [4]
RP Nucleic Acids Res. 11:2733-2744(1983).
CC -1- FUNCTION: COLLAGEN TYPE III OCCURS IN MOST SOFT CONNECTIVE TISSUES
CC -1- ALONG WITH TYPE I COLLAGEN.
CC -1- SUBUNIT: TRIMERS OF IDENTICAL ALPHA 1(III) CHAINS. THE CHAINS ARE
CC LINKED TO EACH OTHER BY INTERCHAIN DISULFIDE BONDS. TRIMERS ARE
CC ALSO CROSS-LINKED VIA HYDROXYLYSINES.
CC -1- PTM: Prolines at the third position of the tripeptide repeating
CC unit (G-X-Y) are hydroxylated in some or all of the chains.
CC -1- SIMILARITY: Contains 1 WFC domain.
CC -----
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DR EMBL: U07973; AAA83407.1; -
DR EMBL: X00822; CAB52686.1; -
DR EMBL: X00823; CAB52686.1; JOINED.
DR EMBL: X00826; CA253397.1; ALT_SEQ.
DR EMBL: X00825; CA253397.1; JOINED.
DR EMBL: X00827; CA253398.1; -
DR EMBL: X00828; CA253399.1; -
DR EMBL: X00830; CA25401.1; -
DR EMBL: X00831; CA25402.1; -
DR EMBL: X02302; AAD15289.1; -
DR EMBL: K02301; AAD15289.1; -
DR EMBL: M36662; AAA18519.1; ALT_SEQ.
DR PIR: A05269; A05269.
DR PIR: I50694; I50694.
DR InterPro: IPR000087; Collagen.
DR InterPro: IPR000885; Fib.collagen.C.
DR InterPro: IPR001007; WFC.
DR ProDom: PD000007; C1g_helix_1.
DR ProDom: PD002078; Fib_collagen_C_1.
DR SMART: SM00338; COLF1_1.
DR SMART: SM00214; WVC_1.
DR PROSITE: PS01208; WFC_1_1.
DR PROSITE: PS50184; WFC_2_1.
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KW Glycoprotein; Collagen; Signal.
FT SIGNAL 1 23
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FT CHAIN 145 1003
FT PROPEP 1004 1262
FT FT
FT FT
FT DOMAIN 145 164
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FT DOMAIN 165 994
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FT FT
FT DOMAIN 995 1003
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FT DISULFID 994 994
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FT DISULFID 995 995
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FT MOD_RES 859 859
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FT FT
FT CONFLICT 96 96
FT CONFLICT 1132 1132
SQ SEQUENCE 1262 AA; 121249 MW; 96ABE7B2E9DEB43D CRC64;

Query Match 71.2%; Score 57; DB 1; Length 1262;
Best Local Similarity 69.2%; Pred. No. 0.56;
Matches 9; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 GTPGPGIAGGRC 13
Db 851 GPPGPGVKGRC 863

RESULT 15
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AC P08121; O61429; Q9CRN7;
DT 01-AUG-1988 (Rel. 08, Created)
DT 15-JUL-1999 (Rel. 38, Last sequence update)
DT 15-SEP-2003 (Rel. 42, Last annotation update)
DE Collagen alpha 1(III) chain precursor.
GN COL3A1.
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
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RP SEQUENCE FROM N.A.

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RC STRAIN=C57BL/6 X DBA; TISSUE=Embryo;  
 RX MEDLINE=95011609; PubMed=7926795;  
 RA Toman D., de Crombrughe B.;  
 RT "The mouse type-III procollagen-encoding gene: genomic cloning and  
 complete DNA sequence."  
 RL Gene 147:161-168(1994).  
 RN (2)  
 RP SEQUENCE OF 1-488 FROM N.A.  
 RX MEDLINE=88167858; PubMed=3443309;  
 RA Wood L., Thériault N., Vogel G.;  
 RT "Complete nucleotide sequence of the N-terminal domains of the murine  
 alpha-1(I)-type-III collagen chain."  
 RL Gene 61:225-230(1987).  
 RN (3)  
 RP SEQUENCE OF 1-28 FROM N.A.  
 RX MEDLINE=85131189; PubMed=3972847;  
 RA Liu G., Mudryj M., de Crombrughe B.;  
 RT "Identification of the promoter and first exon of the mouse alpha 1  
 (III) collagen gene."  
 RL J. Biol. Chem. 260:3773-3777(1985).  
 RN (4)  
 RP SEQUENCE OF 810-1464 FROM N.A.  
 RX STRAIN=C57BL/6J; TISSUE=Embryonic head;  
 MEDLINE=21083660; PubMed=11217851;  
 RA Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,  
 Alakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,  
 Akawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamana K. I.,  
 Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,  
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 Flisbachman W., Gaasterland T., Gissi C., King B., Kochwa H.,  
 Kuhl P., Lewis S., Matsuo Y., Nakaido I., Pesole G., Quackenbush J.,  
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 Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,  
 Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K. F.,  
 Suzuki H., Toyooka K., Wang K. H., Weitz C., Whitaker C., Wilming L.,  
 Wyszewski-Boris A., Yoshida K., Hasegawa Y., Kawai H., Kohlsuki S.,  
 Hayashizaki Y.;  
 RA "Functional annotation of a full-length mouse cDNA collection."  
 RT Nature 409:685-690(2001).  
 RL (5)  
 RP SEQUENCE OF 1442-1464 FROM N.A.  
 RX STRAIN=C57BL/6;  
 MEDLINE=91274355; PubMed=2054384;  
 RA Metsaeranta M., Toman D., de Crombrughe B., Vuorio E.;  
 RT "Specific hybridization probes for mouse type I, II, III and IX  
 collagen mRNAs."  
 RL Biochim. Biophys. Acta 1089:241-243(1991).  
 CC -1- FUNCTION: COLLAGEN TYPE III OCCURS IN MOST SOFT CONNECTIVE TISSUES  
 CC ALONG WITH TYPE I COLLAGEN.  
 CC -1- SUBUNIT: TRIMERS OF IDENTICAL ALPHA 1(III) CHAINS. THE CHAINS ARE  
 CC LINKED TO EACH OTHER BY INTERCHAIN DISULFIDE BONDS. TRIMERS ARE  
 CC ALSO CROSS-LINKED VIA HYDROXYLYSINES.  
 CC -1- PTM: Proline residues at the third position of the tripeptide  
 CC repeating unit (G-X-Y) are hydroxylated in some or all of the  
 CC chains.  
 CC -1- PTM: O-linked glycan consists of a Glc-Gal disaccharide bound to  
 CC the oxygen atom of a post-translationally added hydroxyl group (By  
 CC similarity).  
 CC -1- SIMILARITY: Contains 1 WMEC domain.  
 CC  
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DR EMBL: X52046; CAA36279.1; -;  
 DR EMBL: M18933; AAA37338.1; -;  
 DR EMBL: K03037; -; NOT\_ANNOTATED\_CDS.  
 DR EMBL: AK019448; BAB31724.1; -;  
 DR EMBL: X57983; CAA41048.1; -;  
 DR PIR: A27353; A27353.  
 DR PIR: S59856; S59856.  
 DR MGI: 88453; Col3a1.  
 DR InterPro: IPR00087; Collagen.  
 DR InterPro: IPR00085; Fib\_collagen\_C.  
 DR InterPro: IPR001007; WVC\_C.  
 DR Pfam: PF01410; COLFI; 1.  
 DR Pfam: PF01391; Collagen; 18.  
 DR Prodom: PP000007; Ctg\_helix; 1.  
 DR Prodom: PP002078; Fib\_collagen\_C; 1.  
 DR SMART: SM00038; COLFI; 1.  
 DR SMART: SM00214; WVC; 1.  
 DR PROSITE: PS01208; WVC\_1; 1.  
 DR PROSITE: PS0184; WVC\_2; 1.  
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 KW Glycoprotein; Collagen; signal.  
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 FT PROPEP 1204 1464 CARBOXYL-TERMINAL PROPEPTIDE.  
 FT DOMAIN 31 90 WVC.  
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 FT MOD\_RES 859 859 HYDROXYLATION (BY SIMILARITY).  
 FT MOD\_RES 976 976 HYDROXYLATION (BY SIMILARITY).  
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 FT DISULFD 1196 1196 INTERCHAIN (BY SIMILARITY).  
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Query Match 71.2%; Score 57; DB 1; Length 1464;  
 Best Local Similarity 69.2%; Pred. No. 0.65;  
 Matches 9; Conservative 2; Mismatches 2; Indels 0; Gaps 0;  
 Oy 1 GTPGPGIAGORG 13  
 Db 851 GTPGPGVKGGRG 863

Search completed: August 29, 2003, 18:27:05  
 Job time : 24 secs



GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: August 29, 2003, 18:21:45; Search time 96 Seconds  
(without alignments)  
40.321 Million cell updates/sec

Title: US-09-935-417-1  
Perfect score: 80  
Sequence: 1 GTPGPGIAGRGV 15

Scoring table: BLOSUM62  
Gapop 10.0, Gapext 0.5

Searched: 830525 seqs, 258052604 residues

Total number of hits satisfying chosen parameters: 830525

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database :

SPTREMBL\_23:\*

- 1: sp\_archaea:\*
- 2: sp\_bacteria:\*
- 3: sp\_fungi:\*
- 4: sp\_human:\*
- 5: sp\_invertebrate:\*
- 6: sp\_mammal:\*
- 7: sp\_mpc:\*
- 8: sp\_organelle:\*
- 9: sp\_phage:\*
- 10: sp\_plant:\*
- 11: sp\_protent:\*
- 12: sp\_virus:\*
- 13: sp\_unclassified:\*
- 14: sp\_virus:\*
- 15: sp\_bacteriophage:\*
- 17: sp\_archaeal:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	80	100.0	230	11	Q9R149
2	80	100.0	589	11	Q991L6
3	80	100.0	1453	11	Q63079
4	80	100.0	1461	4	Q76045
5	80	100.0	1464	4	Q8N473
6	75	93.8	809	13	Q93485
7	75	93.8	1449	13	Q910C0
8	72	92.5	1450	13	Q9Y1B4
9	72	90.0	678	13	Q93486
10	72	90.0	1458	13	Q910B9
11	71	88.8	113	11	Q8V172
12	71	88.8	347	6	Q9XT25
13	71	88.8	826	11	Q8K0M6
14	71	88.8	886	11	Q8CEP7
15	71	88.8	1160	4	Q14046
16	71	88.8	1418	6	Q28396

17	71	88.8	1418	13	Q9W7R9	Q9W7R9 cynops pyr
18	71	88.8	1419	13	Q63123	Q63123 rattus norv
19	71	88.8	1420	11	Q90W37	Q90W37 gallus gall
20	71	88.8	1442	11	Q62031	Q62031 mus musculu
21	71	88.8	1442	11	Q62033	Q62033 mus musculu
22	71	88.8	1459	11	Q62032	Q62032 mus musculu
23	71	88.8	1487	4	Q14047	Q14047 homo sapien
24	71	88.8	1487	6	Q77753	Q77753 canis famill
25	68	85.0	1486	13	Q91717	Q91717 xenopus lae
26	64	80.0	1445	13	Q93251	Q93251 rana catesb
27	64	80.0	1447	13	Q91B91	Q91B91 xenopus lae
28	63	78.8	441	4	Q96A83	Q96A83 homo sapien
29	60	75.0	100	4	Q9Y3P3	Q9Y3P3 homo sapien
30	60	75.0	441	4	Q96A84	Q96A84 homo sapien
31	58	72.5	290	5	Q26054	Q26054 paracentrot
32	58	72.5	1258	13	Q8AW11	Q8AW11 brachydantio
33	58	72.5	1347	4	Q960B3	Q960B3 homo sapien
34	58	72.5	1464	11	Q8BKX2	Q8BKX2 mus musculu
35	58	72.5	1497	11	Q61431	Q61431 mus musculu
36	57	71.2	310	13	Q90612	Q90612 gallus gall
37	57	71.2	832	4	Q961F7	Q961F7 homo sapien
38	57	71.2	998	11	Q8CFM4	Q8CFM4 mus musculu
39	57	71.2	1222	11	Q8K173	Q8K173 mus musculu
40	57	71.2	1414	5	Q26634	Q26634 strongyloce
41	57	71.2	1464	11	Q8BLW4	Q8BLW4 mus musculu
42	57	71.2	1860	4	Q81ZC6	Q81ZC6 homo sapien
43	56	70.0	690	13	Q81G18	Q81G18 brachydantio
44	56	70.0	1366	4	Q15177	Q15177 homo sapien
45	56	70.0	1621	4	Q9H4R9	Q9H4R9 homo sapien

#### ALIGNMENTS

RESULT 1	Q9R149	PRELIMINARY:	PRT:	230 AA.
AC	Q9R149:			
DT	01-MAY-2000 (TREMBLrel. 13, Created)			
DT	01-MAY-2000 (TREMBLrel. 13, Last sequence update)			
DE	01-OCT-2002 (TREMBLrel. 22, Last annotation update)			
DE	Pro-alpha-1 type 1 collagen (Fragment).			
OC	Cavia porcellus (Guinea pig).			
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;			
OC	Mammalia; Eutheria; Rodentia; Hystriocognathi; Cavidae; Cavia.			
OX	NCBI_TaxID=10141;			
RN	[1]			
RP	SEQUENCE FROM N.A.			
RC	STRAIN=Hartley; TISSUE=Trachea;			
RA	Morishima Y., Uchida Y., Nomura A., Ishii Y., Sakamoto T.,			
RA	Sekizawa K.;			
RT	"Guinea-pig pro-alpha-1 type 1 collagen expression in injured tracheal			
RT	epithelium.";			
RT	Submitted (JUL-1999) to the EMBL/GenBank/DBJ databases.			
DR	EMBL: AF169346; AAD49346.1; -			
DR	InterPro: IPR000087; Collagen.			
DR	PIfam: PF01391; Collagen; 4.			
KW	Collagen.			
FT	NON_TER	1	230	1
FT	NON_TER	230	230	1
SQ	SEQUENCE	230 AA:	20425 MW;	1AA65F92779D9A71 CRC64;
Query Match				
Best Local Similarity	100.0%;	Score 80;	DB 11;	Length 230;
Matches	15; Conservative	0;	Pred. No. 0.00015;	
		0;	Mismatches	0;
			Indels	0;
			Gaps	0;
OY	1 GTPGPGIAGRGV 15			
Db				
	78 GTPGPGIAGRGV 92			
RESULT 2	Q991L6			

ID 099L6 PRELIMINARY; PRT; 589 AA.  
 AC 099L6;  
 DT 01-JUN-2001 (TREMBLrel. 17, Created)  
 DT 01-JUN-2001 (TREMBLrel. 17, Last sequence update)  
 DT 01-OCT-2002 (TREMBLrel. 22, Last annotation update)  
 DE Hypothetical 58.8 kDa protein (Fragment).  
 GN COL1A1.  
 OS Mus musculus (Mouse).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
 OX NCBI\_TaxID=10090;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RA Strausberg R.;  
 RL Submitted (FEB-2001) to the EMBL/GenBank/DBJ databases.  
 DR EMBL; BC003198; AA03198.1;  
 DR MGD; MGI:88467; Collal.  
 DR InterPro; IPR000885; Fib\_collagen.  
 DR InterPro; IPR000885; Fib\_collagen\_C.  
 DR Pfam; PF01410; COLF1; 1.  
 DR ProDom; PD002078; Fib\_collagen\_C; 1.  
 DR SMART; SM00038; COLF1; 1.  
 DR Hypothetical protein.  
 KW NON\_TER  
 FT SEQUENCE 589 AA; 58805 MW; 81847495SE05CEP CRC64;

Query Match 100.0%; Score 80; DB 11; Length 589;  
 Best Local Similarity 100.0%; Pred. No. 0.00039;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GTPPGGIAGGRCVV 15  
 Db 72 GTPPGGIAGGRCVV 86

## RESULT 3

ID 063079 PRELIMINARY; PRT; 1453 AA.  
 AC 063079;  
 DT 01-NOV-1996 (TREMBLrel. 01, Created)  
 DT 01-JUN-1998 (TREMBLrel. 06, Last sequence update)  
 DT 01-OCT-2002 (TREMBLrel. 22, Last annotation update)  
 DE Collagen alpha1 (Fragment).  
 OS Rattus norvegicus (Rat).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.  
 OX NCBI\_TaxID=10116;  
 RN [1]  
 RP SEQUENCE OF 1-1092 FROM N.A.  
 RC STRAIN-Sprague-Dawley; TISSUE-Tooth;  
 RA Brandsten C., Lundmark C., Christerson C., Hammarstrom L., Murtz T.;  
 RT "Expression of Collagen alpha1(I) mRNA variants during Tooth and Bone  
 Formation in the Rat."  
 RT J. Dent. Res. 0:0-0(0).  
 RL [2]  
 RN SEQUENCE FROM N.A.  
 RC STRAIN-Sprague-Dawley; TISSUE-Tooth;  
 RA Murtz T.;  
 RL Submitted (FEB-1998) to the EMBL/GenBank/DBJ databases.  
 DR EMBL; Z78279; CAB01633.1;  
 DR InterPro; IPR000885; Collagen.  
 DR InterPro; IPR001007; VWF\_C.  
 DR Pfam; PF01410; COLF1; 1.  
 DR Pfam; PF01391; Collagen; 18.  
 DR ProDom; PD000007; Collagen; 3.  
 DR ProDom; PD002078; Fib\_collagen\_C; 1.  
 DR SMART; SM00038; COLF1; 1.  
 DR SMART; SM00214; WVC1; 1.  
 DR PROSITE; PS01208; WVC1; 1.  
 KW Collagen.  
 FT NON\_TER 1 1

SQ SEQUENCE 1453 AA; 137887 MW; E6896BDC19A4A1D8 CRC64;

Query Match 100.0%; Score 80; DB 11; Length 1453;  
 Best Local Similarity 100.0%; Pred. No. 0.00096;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GTPPGGIAGGRCVV 15  
 Db 936 GTPPGGIAGGRCVV 950

## RESULT 4

ID 076045 PRELIMINARY; PRT; 1461 AA.  
 AC 076045;  
 DT 01-NOV-1998 (TREMBLrel. 08, Created)  
 DT 01-NOV-1998 (TREMBLrel. 12, Last sequence update)  
 DT 01-OCT-2002 (TREMBLrel. 22, Last annotation update)  
 DE Pro alpha 1(I) collagen.  
 GN COL1A1.  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.  
 OX NCBI\_TaxID=9606;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RA Chu M.L., de Wet W., Bernard M., Ramirez F.;  
 RT "Fine structural analysis of the human pro-alpha 1 (I) collagen gene.  
 RT promoter structure, Alu repeats, and polymorphic transcripts."  
 RT J. Biol. Chem. 260:2315-2320(1985).  
 RN [2]  
 RP SEQUENCE FROM N.A.  
 RA MEDLINE=88329734; PubMed=2843432;  
 RX D'Alessio M., Bernard M., Pretorius P.J., de Wet W., Ramirez F.;  
 RT "Complete nucleotide sequence of the region encompassing the first  
 RT twenty-five exons of the human pro alpha 1(I) collagen gene."  
 RL Gene 67:105-115(1988).  
 RN [3]  
 RP SEQUENCE FROM N.A.  
 RA MEDLINE=89025644; PubMed=3178743;  
 RX Tromp G., Kulvanliem H., Stacey A., Shikata H., Baldwin C.T.,  
 RA Jaenisch R., Prockop D.J.;  
 RT "Structure of a full-length cDNA clone for the prepro alpha 1(I) chain  
 RT of human type I procollagen."  
 RL Biochem. J. 253:919-922(1988).  
 RN [4]  
 RP SEQUENCE FROM N.A.  
 RX MEDLINE=91138770; PubMed=1995349;  
 RA Maatta A., Bornstein P., Penttinen R.P.;  
 RT "Highly conserved sequences in the 3'-untranslated region of the  
 RT COL1A1 gene bind cell-specific nuclear proteins."  
 RL FEBS Lett. 279:9-13(1991).  
 RN [5]  
 RP SEQUENCE FROM N.A.  
 RX MEDLINE=92157916; PubMed=1787829;  
 RA Westerhausen A., Constantinou C.D., Pack M., Peng M.Z., Hanning C.,  
 RA Olsen A.S., Prockop D.J.;  
 RT "Completion of the last half of the structure of the human gene for  
 RT the pro alpha 1 (I) chain of type I procollagen (COL1A1)."  
 RL Matrix 11:375-379(1991).  
 RN [6]  
 RP SEQUENCE FROM N.A.  
 RX MEDLINE=98107942; PubMed=9443882;  
 RA Korhonen J., Ala-Korhonen L., De Paeppe A., Nuytink L., Barley J.,  
 RA Prockop D.J.;  
 RT "Analysis of the COL1A1 and COL1A2 genes by PCR amplification and  
 RT scanning by conformation-sensitive gel electrophoresis identifies only  
 RT COL1A1 mutations in 15 patients with osteogenesis imperfecta type I;  
 RT identification of common sequences of null-allele mutations."  
 RL Am. J. Hum. Genet. 62:98-110(1998).  
 RN [7]  
 RP SEQUENCE FROM N.A.

RA Korkko J.M., Earley J.J., Nuytink L., Depaepe A., Prockop D.J.,  
 RA Ala-Korkko L.;  
 RL Submitted (May-1999) to the EMBL/Genbank/DBJ databases.  
 DR EMBL, AF017178; AAB94054.2; -;  
 DR InterPro: IPR000087; Collagen.  
 DR InterPro: IPR000885; Fib\_collagen\_C.  
 DR InterPro: IPR001007; VWF\_C.  
 DR Pfam: PF01410; COLFI; 1.  
 DR Pfam: PF01391; Collagen; 18.  
 DR Pfam: PF00093; VWC; 1.  
 DR ProDom: PD000007; Collagen; 2.  
 DR ProDom: PD002078; Fib\_collagen\_C; 1.  
 DR SMART: SM00038; COLFI; 1.  
 DR SMART: SM00214; VWC; 1.  
 DR PROSITE: PS01208; VWF\_C; 1.  
 DR Collagen.  
 KW COLLAGEN.  
 SQ SEQUENCE 1461 AA; 138630 MW; 9ACF6DE30EA78E21 CRC64;  
 Query Match 100.0%; Score 80; DB 4; Length 1461;  
 Best Local Similarity 100.0%; Pred. No. 0.00097;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 GTPPGGIAGGCGV 15  
 DB 944 GTPPGGIAGGCGV 958  
 RESULT 5  
 Q8N473 PRELIMINARY; PRT; 1464 AA.  
 AC Q8N473;  
 DT 01-OCT-2002 (TREMBLrel. 22, Created)  
 DT 01-OCT-2002 (TREMBLrel. 22, Last sequence update)  
 DT 01-MAR-2003 (TREMBLrel. 23, Last annotation update)  
 DE Hypothetical protein.  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 OX NCBI\_TaxID=9606;  
 OX 11;  
 RN SEQUENCE FROM N.A.  
 RP TISSUE=Brain;  
 RC Strausberg R.;  
 RL Submitted (Aug-2002) to the EMBL/Genbank/DBJ databases.  
 DR EMBL, BC036531; AAH36531.1; -;  
 DR InterPro: IPR000087; Collagen.  
 DR InterPro: IPR000885; Fib\_collagen\_C.  
 DR InterPro: IPR001007; VWF\_C.  
 DR Pfam: PF01410; COLFI; 1.  
 DR Pfam: PF01391; Collagen; 18.  
 DR ProDom: PD000007; Collagen; 2.  
 DR ProDom: PD002078; Fib\_collagen\_C; 1.  
 DR SMART: SM00038; COLFI; 1.  
 DR SMART: SM00214; VWC; 1.  
 DR PROSITE: PS01208; VWF\_C; 1.  
 DR Hypothetical protein: Collagen.  
 KW COLLAGEN.  
 SQ SEQUENCE 1464 AA; 139011 MW; B0581FBD1C99DDE8 CRC64;  
 Query Match 100.0%; Score 80; DB 4; Length 1464;  
 Best Local Similarity 100.0%; Pred. No. 0.00097;  
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 GTPPGGIAGGCGV 15  
 DB 947 GTPPGGIAGGCGV 961  
 RESULT 6  
 O93485 PRELIMINARY; PRT; 809 AA.  
 AC O93485;  
 DT 01-NOV-1998 (TREMBLrel. 08, Created)  
 DT 01-NOV-1998 (TREMBLrel. 08, Last sequence update)

DT 01-OCT-2002 (TREMBLrel. 22, Last annotation update)  
 DE Alpha 1 type I collagen (Fragment).  
 OS Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Actinopterygii; Neopterygii; Teleostei; Euteleostei;  
 OC Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.  
 OX NCBI\_TaxID=8022;  
 RN 11;  
 RP SEQUENCE FROM N.A.  
 RC TISSUE=Fibroblast;  
 RA Saito M., Kunisaki N., Hirono I., Aoki T., Ishida M., Urano N.,  
 RA Kimura S.;  
 RT "Partial characterization of cDNA clones encoding the three distinct  
 RT pro alpha chains of type I collagen from rainbow trout.";  
 RL Fisheries Sci. 64:780-786(1998).  
 DR EMBL, AB008373; BAA3380.1; -;  
 DR InterPro: IPR000087; Collagen.  
 DR InterPro: IPR000885; Fib\_collagen\_C.  
 DR Pfam: PF01391; Collagen; 9.  
 DR ProDom: PD000007; Collagen; 1.  
 DR ProDom: PD002078; Fib\_collagen\_C; 1.  
 DR SMART: SM00038; COLFI; 1.  
 DR Collagen.  
 KW NON\_TER.  
 FT  
 SQ SEQUENCE 809 AA; 78164 MW; 68C056A7640FCAB1 CRC64;  
 Query Match 93.8%; Score 75; DB 13; Length 809;  
 Best Local Similarity 86.7%; Pred. No. 0.0031;  
 Matches 13; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
 QY 1 GTPPGGIAGGCGV 15  
 DB 292 GTPPGGIAGGCGV 306  
 RESULT 7  
 Q910C0 PRELIMINARY; PRT; 1449 AA.  
 AC Q910C0;  
 DT 01-DEC-2001 (TREMBLrel. 19, Created)  
 DT 01-DEC-2001 (TREMBLrel. 19, Last sequence update)  
 DT 01-MAR-2003 (TREMBLrel. 23, Last annotation update)  
 DE Collagen a1(I).  
 GN COL1A1.  
 OS Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Actinopterygii; Neopterygii; Teleostei; Euteleostei;  
 OC Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.  
 OX NCBI\_TaxID=8022;  
 OX 11;  
 RN SEQUENCE FROM N.A.  
 RP MEDLINE=21257802; PubMed=11358497;  
 RA Saito M., Takenouchi Y., Kunisaki N., Kimura S.;  
 RT "Complete primary structure of rainbow trout type I collagen  
 RT consisting of alpha1(I)2(1)3(I) heterotrimer.";  
 RL Eur. J. Biochem. 268:2817-2827(2001).  
 DR EMBL, AB052835; BAB55661.1; -;  
 DR InterPro: IPR000087; Collagen.  
 DR InterPro: IPR000885; Fib\_collagen\_C.  
 DR InterPro: IPR001007; VWF\_C.  
 DR Pfam: PF01410; COLFI; 1.  
 DR Pfam: PF01391; Collagen; 18.  
 DR Pfam: PF00093; VWC; 1.  
 DR ProDom: PD000007; Collagen; 2.  
 DR ProDom: PD002078; Fib\_collagen\_C; 1.  
 DR SMART: SM00038; COLFI; 1.  
 DR SMART: SM00214; VWC; 1.  
 DR PROSITE: PS01208; VWF\_C; 1.  
 DR Collagen.  
 KW COLLAGEN.  
 SQ SEQUENCE 1449 AA; 137117 MW; 62EEF8A7BFD652B8 CRC64;  
 Query Match 93.8%; Score 75; DB 13; Length 1449;

DE Collagen type II (Fragment).  
OS *Cavia porcellus* (Guinea pig).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

```

OC Mammalia; Eutheria; Rodentia; Hystriognathi; Caviidae; Cavia.
OX NCBI_TaxId=10141;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=Hartley;
RA Huebner J.L., Clark A.E., Kraus V.B., Otterness I.G.;
RT "Collagen type II in the guinea pig.";
RL Submitted (May-1999) to the EMBL/GenBank/DBJ databases.
DR EMBL: AF152862; AAL55558.1; -
DR InterPro: IPR000087; Collagen.
DR Pfam: PF01391; Collagen.1.
DR ProDom: PD000007; Collagen; 1.
KW Collagen.
FT NON_TER
FT NON_TER
SQ SEQUENCE 113 AA; 10284 MW; F7861901127A9BCE CRC64;

Query Match
Best Local Similarity 88.8%; Score 71; DB 11; Length 113;
Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGV 15
DB 89 GPPGPGIAGRGIV 103

RESULT 12
Q9XT25 PRELIMINARY; PRT; 347 AA.
AC Q9XT25;
DT 01-NOV-1999 (TREMBlrel. 12, Created)
DT 01-NOV-1999 (TREMBlrel. 12, Last sequence update)
DT 01-OCT-2002 (TREMBlrel. 22, Last annotation update)
DE Type II collagen cyanogen bromide fragment CB10 (Fragment).
OS Bos taurus (Bovine).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidea;
OC Bovidae; Bovinae; Bos.
OX NCBI_TaxId=9913;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Cartilage;
RA Tang B., Chiang T.M., Brand D.D., Gumanovskaya M.L., Stuart J.M.,
RA Kang A.H., Myers L.K.;
RT "Molecular Definition and Characterization of Recombinant Bovine CB8
RT And CB10: Immunogenicity and Arthritisogenicity.";
RL J. Clin. Immunol. 0:0-0(1999).
DR EMBL: AF138883; AAD42346.1; -
DR InterPro: IPR000087; Collagen.
DR Pfam: PF01391; Collagen; 6.
DR ProDom: PD000007; Collagen; 2.
KW Collagen.
FT NON_TER
FT NON_TER
SQ SEQUENCE 347 AA; 31085 MW; 5D41COAF34089DF6 CRC64;

Query Match
Best Local Similarity 88.8%; Score 71; DB 6; Length 347;
Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGV 15
DB 219 GPPGPGIAGRGIV 233

RESULT 13
Q8K0N6 PRELIMINARY; PRT; 826 AA.
AC Q8K0N6;
DT 01-OCT-2002 (TREMBlrel. 22, Created)
DT 01-OCT-2002 (TREMBlrel. 22, Last sequence update)
DT 01-OCT-2002 (TREMBlrel. 22, Last annotation update)
DT 01-MAR-2003 (TREMBlrel. 23, Last annotation update)
DE Similar to procollagen, type II, alpha 1 (Fragment).

```

```

OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxId=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Retina;
RA Strusberg R.;
RL Submitted (JUN-2002) to the EMBL/GenBank/DBJ databases.
DR EMBL: BC030913; AAH30913.1; -
DR InterPro: IPR000087; Collagen.
DR InterPro: IPR000885; Fib_collagen_C.
DR Pfam: PF01410; COLFI.1.
DR Pfam: PF01391; Collagen; 9.
DR ProDom: PD000007; Collagen; 2.
DR ProDom: PD002078; Fib_collagen_C; 1.
DR SMART: SM00038; COLFI; 1.
KW Collagen.
FT NON_TER
FT NON_TER
SQ SEQUENCE 826 AA; 80124 MW; B5BD721772BDAF24 CRC64;

Query Match
Best Local Similarity 88.8%; Score 71; DB 11; Length 826;
Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGV 15
DB 308 GPPGPGIAGRGIV 322

RESULT 14
O8CEF7 PRELIMINARY; PRT; 886 AA.
AC O8CEF7;
DT 01-MAR-2003 (TREMBlrel. 23, Created)
DT 01-MAR-2003 (TREMBlrel. 23, Last sequence update)
DT 01-MAR-2003 (TREMBlrel. 23, Last annotation update)
DE Procollagen.
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxId=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Head;
RA MEDLINE=22354683; PubMed=12466651;
RA The FANTOM Consortium.
RA The RIKEN Genome Exploration Research Group Phase I & II Team;
RT "Analysis of the mouse transcriptome based on functional annotation of
RT 60,770 full-length cDNAs.";
RL Nature 420:563-573(2002).
DR EMBL: AK028295; BAC25865.1; -
DR Nature 420:563-573(2002).
SQ SEQUENCE 886 AA; 85536 MW; 47A70AA0DBBF4F45 CRC64;

Query Match
Best Local Similarity 88.8%; Score 71; DB 11; Length 886;
Matches 12; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 GTPGPGIAGRGV 15
DB 368 GPPGPGIAGRGIV 382

RESULT 15
Q14046 PRELIMINARY; PRT; 1160 AA.
AC Q14046;
DT 01-NOV-1996 (TREMBlrel. 01, Created)
DT 01-NOV-1996 (TREMBlrel. 01, Last sequence update)
DT 01-OCT-2002 (TREMBlrel. 22, Last annotation update)
DE COL2A1 protein precursor (Fragment).
GN COL2A1
OS Homo sapiens (Human).

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